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(54) **Luciferase gene and novel recombinant DNA as well as a method for production of luciferase.**

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Description

BACKGROUND OF THE INVENTION

1. Field of the Invention

This invention relates to a luciferase gene derived from *Luciola lateralis* (HEIKE firefly) and a novel recombinant DNA having integrated the gene therein as well as a method of producing a luciferase using the recombinant DNA.

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2. Description of the Prior Art

Luciferase from fireflies belonging to the genus *Luciola* is merely obtained by isolating and purifying from the collected fireflies belonging to the genus *Luciola* [Proc. Natl. Acad. Sci., 74 (7), 2799-2802 (1977)].

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Luciferases are very effectively usable, e.g., for quantitative determination of ATP.

Since luciferases described above are derived from insects, however, fireflies belonging to the genus *Luciola* must be collected from the natural world to produce luciferases; alternatively, such fireflies must be cultivated and luciferases should be isolated and refined from the fireflies so that much time and labors are required for the production.

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As a result of various investigations to solve the foregoing problems, the present inventors have found that by producing a recombinant DNA by inserting DNA containing a *Luciola lateralis*-derived luciferase-coding gene into a vector DNA and culturing in a medium a luciferase-producing microorganism belonging to the genus *Escherichia* and bearing the recombinant DNA, luciferase can be efficiently produced in a short period of time. As a result of further investigations on luciferase gene derived from *Luciola lateralis*, the present inventors have also succeeded in isolating a luciferase gene derived from *Luciola lateralis* and determining its structure, for the first time. This invention has thus been accomplished.

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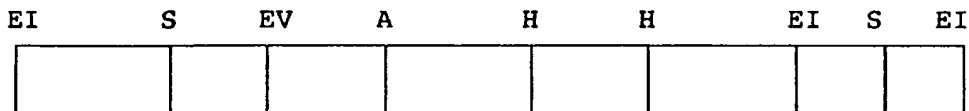
SUMMARY OF THE INVENTION

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According to this invention, there are provided:

(1) A *Luciola lateralis*-derived luciferase gene defined by a restriction enzyme map described below:

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wherein EI represents Eco RI, S represents Ssp I, EV represents Eco RV, A represents Apa I and H represents Hpa I.

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(2) A luciferase gene according to (1) which encodes an amino acid sequence shown below:

										10
5	Met	Glu	Asn	Met	Glu	Asn	Asp	Glu	Asn	Ile
										20
	Val	Tyr	Gly	Pro	Glu	Pro	Phe	Tyr	Pro	Ile
										30
10	Glu	Glu	Gly	Ser	Ala	Gly	Ala	Gln	Leu	Arg
										40
	Lys	Tyr	Met	Asp	Arg	Tyr	Ala	Lys	Leu	Gly
										50
15	Ala	Ile	Ala	Phe	Thr	Asn	Ala	Leu	Thr	Gly
										60
	Val	Asp	Tyr	Thr	Tyr	Ala	Glu	Tyr	Leu	Glu
										70
20	Lys	Ser	Cys	Cys	Leu	Gly	Glu	Ala	Leu	Lys
										80
	Asn	Tyr	Gly	Leu	Val	Val	Asp	Gly	Arg	Ile
										90
25	Ala	Leu	Cys	Ser	Glu	Asn	Cys	Glu	Glu	Phe
										100
	Phe	Ile	Pro	Val	Leu	Ala	Gly	Leu	Phe	Ile
										110
30	Gly	Val	Gly	Val	Ala	Pro	Thr	Asn	Glu	Ile
										120
	Tyr	Thr	Leu	Arg	Glu	Leu	Val	His	Ser	Leu
										130
35	Gly	Ile	Ser	Lys	Pro	Thr	Ile	Val	Phe	Ser
										140
	Ser	Lys	Lys	Gly	Leu	Asp	Lys	Val	Ile	Thr
										150
40	Val	Gln	Lys	Thr	Val	Thr	Ala	Ile	Lys	Thr
										160
	Ile	Val	Ile	Leu	Asp	Ser	Lys	Val	Asp	Tyr
										170
45	Arg	Gly	Tyr	Gln	Ser	Met	Asp	Asn	Phe	Ile
										180
	Lys	Lys	Asn	Thr	Pro	Gln	Gly	Phe	Lys	Gly
										190
50	Ser	Ser	Phe	Lys	Thr	Val	Glu	Val	Asn	Arg

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	Lys	Glu	Gln	Val	Ala	Leu	Ile	Met	Asn	Ser	200
	Ser	Gly	Ser	Thr	Gly	Leu	Pro	Lys	Gly	Val	210
5	Gln	Leu	Thr	His	Glu	Asn	Ala	Val	Thr	Arg	220
	Phe	Ser	His	Ala	Arg	Asp	Pro	Ile	Tyr	Gly	230
10	Asn	Gln	Val	Ser	Pro	Gly	Thr	Ala	Ile	Leu	240
	Thr	Val	Val	Pro	Phe	His	His	Gly	Phe	Gly	250
15	Met	Phe	Thr	Thr	Leu	Gly	Tyr	Leu	Thr	Cys	260
	Gly	Phe	Arg	Ile	Val	Met	Leu	Thr	Lys	Phe	270
20	Asp	Glu	Glu	Thr	Phe	Leu	Lys	Thr	Leu	Gln	280
	Asp	Tyr	Lys	Cys	Ser	Ser	Val	Ile	Leu	Val	290
25	Pro	Thr	Leu	Phe	Ala	Ile	Leu	Asn	Arg	Ser	300
	Glu	Leu	Leu	Asp	Lys	Tyr	Asp	Leu	Ser	Asn	310
30	Leu	Val	Glu	Ile	Ala	Ser	Gly	Gly	Ala	Pro	320
	Leu	Ser	Lys	Glu	Ile	Gly	Glu	Ala	Val	Ala	330
35	Arg	Arg	Phe	Asn	Leu	Pro	Gly	Val	Arg	Gln	340
	Gly	Tyr	Gly	Leu	Thr	Glu	Thr	Thr	Ser	Ala	350
40	Ile	Ile	Ile	Thr	Pro	Glu	Gly	Asp	Asp	Lys	360
	Pro	Gly	Ala	Ser	Gly	Lys	Val	Val	Pro	Leu	370
45	Phe	Lys	Ala	Lys	Val	Ile	Asp	Leu	Asp	Thr	380

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	Lys	Lys	Thr	Leu	Gly	Pro	Asn	Arg	Arg	Gly	390
	Glu	Val	Cys	Val	Lys	Gly	Pro	Met	Leu	Met	400
5	Lys	Gly	Tyr	Val	Asp	Asn	Pro	Glu	Ala	Thr	410
	Arg	Glu	Ile	Ile	Asp	Glu	Glu	Gly	Trp	Leu	420
10	His	Thr	Gly	Asp	Ile	Gly	Tyr	Tyr	Asp	Glu	430
	Glu	Lys	His	Phe	Phe	Ile	Val	Asp	Arg	Leu	440
15	Lys	Ser	Leu	Ile	Lys	Tyr	Lys	Gly	Tyr	Gln	450
	Val	Pro	Pro	Ala	Glu	Leu	Glu	Ser	Val	Leu	460
20	Leu	Gln	His	Pro	Asn	Ile	Phe	Asp	Ala	Gly	470
	Val	Ala	Gly	Val	Pro	Asp	Pro	Ile	Ala	Gly	480
25	Glu	Leu	Pro	Gly	Ala	Val	Val	Val	Leu	Glu	490
	Lys	Gly	Lys	Ser	Met	Thr	Glu	Lys	Glu	Val	500
30	Met	Asp	Tyr	Val	Ala	Ser	Gln	Val	Ser	Asn	510
	Ala	Lys	Arg	Leu	Arg	Gly	Gly	Val	Arg	Phe	520
35	Val	Asp	Glu	Val	Pro	Lys	Gly	Leu	Thr	Gly	530
	Lys	Ile	Asp	Gly	Lys	Ala	Ile	Arg	Glu	Ile	540
40	Leu	Lys	Lys	Pro	Val	Ala	Lys	Met			

(3) A luciferase gene according to (1) or (2) which is represented by a nucleotide sequence shown below.

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	ATG	GAA	AAC	ATG	GAG	AAC	GAT	GAA	AAT	ATT	30
5	GTG	TAT	GGT	CCT	GAA	CCA	TTT	TAC	CCT	ATT	60
	GAA	GAG	GGA	TCT	GCT	GGA	GCA	CAA	TTG	CGC	90
10	AAG	TAT	ATG	GAT	CGA	TAT	GCA	AAA	CTT	GGA	120
	GCA	ATT	GCT	TTT	ACT	AAC	GCA	CTT	ACC	GGT	150
15	GTC	GAT	TAT	ACG	TAC	GCC	GAA	TAC	TTA	GAA	180
	AAA	TCA	TGC	TGT	CTA	GGA	GAG	GCT	TTA	AAG	210
20	AAT	TAT	GGT	TTG	GTT	GTT	GAT	GGA	AGA	ATT	240
	GCG	TTA	TGC	AGT	GAA	AAC	TGT	GAA	GAA	TTC	270
25	TTT	ATT	CCT	GTA	TTA	GCC	GGT	TTA	TTT	ATA	300
	GGT	GTC	GGT	GTG	GCT	CCA	ACT	AAT	GAG	ATT	330
30	TAC	ACT	CTA	CGT	GAA	TTG	GTT	CAC	AGT	TTA	360
	GGC	ATC	TCT	AAG	CCA	ACA	ATT	GTA	TTT	AGT	390
35	TCT	AAA	AAA	GGA	TTA	GAT	AAA	GTT	ATA	ACT	420
	GTA	CAA	AAA	ACG	GTA	ACT	GCT	ATT	AAA	ACC	450
40	ATT	GTT	ATA	TTG	GAC	AGC	AAA	GTG	GAT	TAT	480
	AGA	GGT	TAT	CAA	TCC	ATG	GAC	AAC	TTT	ATT	510
45	AAA	AAA	AAC	ACT	CCA	CAA	GGT	TTC	AAA	GGA	540
	TCA	AGT	TTT	AAA	ACT	GTA	GAA	GTT	AAC	CGC	570
50	AAA	GAA	CAA	GTT	GCT	CTT	ATA	ATG	AAC	TCT	600
	TCG	GGT	TCA	ACC	GGT	TTG	CCA	AAA	GGT	GTG	630

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	CAA CTT ACT CAT GAA AAT GCA GTC ACT AGA	660
5	TTT TCT CAC GCT AGA GAT CCA ATT TAT GGA	690
	AAC CAA GTT TCA CCA GGC ACG GCT ATT TTA	720
10	ACT GTA GTA CCA TTC CAT CAT GGT TTT GGT	750
	ATG TTT ACT ACT TTA GGC TAT CTA ACT TGT	780
15	GGT TTT CGT ATT GTC ATG TTA ACG AAA TTT	810
	GAC GAA GAG ACT TTT TTA AAA ACA CTG CAA	840
20	GAT TAC AAA TGT TCA AGC GTT ATT CTT GTA	870
	CCG ACT TTG TTT GCA ATT CTT AAT AGA AGT	900
25	GAA TTA CTC GAT AAA TAT GAT TTA TCA AAT	930
	TTA GTT GAA ATT GCA TCT GGC GGA GCA CCT	960
30	TTA TCT AAA GAA ATT GGT GAA GCT GTT GCT	990
	AGA CGT TTT AAT TTA CCG GGT GTT CGT CAA	1020
35	GGC TAT GGT TTA ACA GAA ACA ACC TCT GCA	1050
	ATT ATT ATC ACA CCG GAA GGC GAT GAT AAA	1080
40	CCA GGT GCT TCT GGC AAA GTT GTG CCA TTA	1110
	TTT AAA GCA AAA GTT ATC GAT CTT GAT ACT	1140
45	AAA AAA ACT TTG GGC CCG AAC AGA CGT GGA	1170
	GAA GTT TGT GTA AAG GGT CCT ATG CTT ATG	1200
50	AAA GGT TAT GTA GAT AAT CCA GAA GCA ACA	1230
	AGA GAA ATC ATA GAT GAA GAA GGT TGG TTG	1260
55	CAC ACA GGA GAT ATT GGG TAT TAC GAT GAA	1290

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      1 3 2 0
GAA AAA CAT TTC TTT ATC GTG GAT CGT TTG
      1 3 5 0
5  AAG TCT TTA ATC AAA TAC AAA GGA TAT CAA
      1 3 8 0
    GTA CCA CCT GCT GAA TTA GAA TCT GTT CTT
      1 4 1 0
10  TTG CAA CAT CCA AAT ATT TTT GAT GCC GGC
      1 4 4 0
    GTT GCT GGC GTT CCA GAT CCT ATA GCT GGT
      1 4 7 0
15  GAG CTT CCG GGA GCT GTT GTT GTA CTT GAA
      1 5 0 0
    AAA GGA AAA TCT ATG ACT GAA AAA GAA GTA
      1 5 3 0
20  ATG GAT TAC GTT GCT AGT CAA GTT TCA AAT
      1 5 6 0
    GCA AAA CGT TTG CGT GGT GGT GTC CGT TTT
      1 5 9 0
25  GTG GAC GAA GTA CCT AAA GGT CTC ACT GGT
      1 6 2 0
    AAA ATT GAC GGT AAA GCA ATT AGA GAA ATA
      1 6 5 0
30  CTG AAG AAA CCA GTT GCT AAG ATG

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According to this invention, there is also provided a novel recombinant DNA obtained by inserting a gene coding for luciferase derived from Luciola lateralis into a vector DNA. According to this invention, there is further provided a method of producing a luciferase which comprises culturing in a medium a microorganism belonging to the genus Escherichia and bearing a recombinant DNA obtained by inserting a gene coding for luciferase derived from Luciola lateralis in a vector DNA, and collecting a luciferase from the culture.

BRIEF DESCRIPTION OF THE DRAWINGS

Fig. 1 shows an optimum pH range of luciferase derived from Luciola lateralis. Fig. 2 shows a stable pH range of luciferase derived from Luciola lateralis. Fig. 3 shows a cleavage map of recombinant plasmid pALf3 DNA with restriction enzymes. Fig. 4 shows a cleavage map of recombinant plasmid pGLf1 DNA with restriction enzymes. Fig. 5 shows a cleavage map of recombinant plasmid pHLf7 DNA with restriction enzymes. Fig. 6 shows a nucleotide sequence of Luciola lateralis-derived luciferase gene in accordance with this invention. Fig. 7 shows an amino acid sequence of polypeptide translated from Luciola lateralis-derived luciferase gene of this invention. Fig. 8 shows a nucleotide sequence of Luciola lateralis-derived luciferase gene and the amino acid sequence corresponding thereto.

DETAILED DESCRIPTION OF THE PREFERRED EMBODIMENTS

Hereafter this invention is described in detail.

In survey of DNA bearing a gene encoding luciferase derived from Luciola lateralis (HEIKE firefly), DNA containing a gene encoding luciferase derived from Luciola cruciata (GENJI firefly) belonging to the same genus, which is one of fireflies, is used as a probe; further in survey of DNA bearing a Luciola cruciata-derived luciferase-coding gene, DNA containing a gene coding for luciferase derived from Photinus pyralis - (American firefly), which is one of fireflies, is used as a probe.

Therefore, production of the gene encoding luciferase from Photinus pyralis is firstly described below. Subsequently, production of the gene encoding luciferase from Luciola cruciata is described and finally, production of the gene encoding luciferase from Luciola lateralis is described.

To prepare m-RNA from the posterior portion of Photinus pyralis which is one of fireflies, m-RNA can be obtained according to methods described in, for example, Molecular Cloning, page 196, Cold Spring Harbor Laboratory (1982), Haruo Ozeki and Reiro Shimura, BUNSHI IDENGAKU JIKKENHO (Experimental Molecular Genetics), pages 66-67 (1983), etc.

Concentration of m-RNA coding for luciferase from the obtained m-RNA can be performed by a method described in, for example, Biomedical Research, 3, 534-540 (1982) or the like.

In this case, anti-luciferase serum to luciferase is used. This serum can be obtained by, for example, Yuichi Yamamura, MEN-EKI KAGAKU (Immunochemistry), pages 43-50 (1973), etc.

Synthesis of c-DNA from the m-RNA coding for luciferase can be performed by methods described in, for example, Mol. Cell Biol., 2, 161 (1982) and Gene, 25, 263 (1983).

Then, the thus obtained c-DNA is integrated into, for example, plasmid pMCE10 DNA {plasmid produced using plasmid pKN305 [plasmid having a promoter of *Escherichia coli* tryptophan operator described in Agr. Biol. Chem., 50, 271 (1986)] and plasmid pMC1843 [plasmid containing *Escherichia coli* β -galactosidase structural gene described in Methods in Enzymology, 100, 293-308 (1983)]}, etc. to produce various recombinant plasmid DNAs. Using these DNAs, transformation of *Escherichia coli* (*E. coli*) DH1 (ATCC 33849), *E. coli* HB101 (ATCC 33694), etc. is effected by the method of Hanahan [DNA Cloning, 1, 109-135 (1985)] to obtain various transformants.

The recombinant plasmid c-DNAs possessed by the thus obtained transformants are plasmids wherein c-DNA has been integrated in the middle of *E. coli* β -galactosidase structural gene. A peptide encoded by c-DNA is expressed as a protein fused with β -galactosidase.

In order to detect c-DNA coding for luciferase from the various transformants described above, the transformants are cultured thereby to express cell protein. By determining if any protein crossing over anti-luciferase serum is present, its detection can be made. Methods described in, for example, Agric. Biol. Chem., 50, 271 (1986) and Anal. Biochem., 112, 195 (1981), etc. can be used for the detection.

Next, after labeling c-DNA of incomplete luciferase with ^{32}P by the nick translation method [Molecular Cloning, pages 109-112, Cold Spring Harbor Laboratory (1982) and J. Mol. Biol., 113, 237-251 (1977)], using the colony hybridization method [Protein, Nucleic Acid & Enzyme, 26, 575-579 (1981)], an *Escherichia coli* strain having plasmid DNA containing Photinus pyralis luciferase c-DNA of 1.8 Kb can be obtained from a Photinus pyralis-derived c-DNA library prepared using plasmid pUC19 DNA (manufactured by Takara Shuzo Co., Ltd.) as a vector.

To obtain a DNA containing the gene coding for luciferase derived from Photinus pyralis from the thus obtained recombinant plasmid DNA, restriction enzymes, e.g., Eco RI and Cla I, are acted on the plasmid DNA at temperatures of 30 to 40°C, preferably at 37°C, for 1 to 24 hours, preferably for 2 hours; the solution obtained after completion of the reaction is subjected to agarose gel electrophoresis [which is described in Molecular Cloning, page 150, Cold Spring Harbor Laboratory (1982)] to obtain the DNA containing the gene coding for luciferase derived from Photinus pyralis.

Next, production of the Luciola cruciata-derived luciferase-coding gene are described below.

Preparation of m-RNA from the posterior portion of Luciola cruciata and synthesis of c-DNA from the m-RNA can be conducted, for example, in quite the same manner as in the preparation of the m-RNA of Photinus pyralis and synthesis of the c-DNA described above.

Then, the thus obtained c-DNA is integrated into a vector DNA, for example, plasmid pUC19 DNA (manufactured by Takara Shuzo Co., Ltd.), etc. to obtain various recombinant plasmid DNAs. Using these DNAs, transformation of *E. coli* DH1 (ATCC 33849), *E. coli* HB101 (ATCC 33694), etc. is effected by the method of Hanahan [DNA Cloning, 1, 109-135 (1985)] to obtain various transformants.

Next, after labeling c-DNA of luciferase derived from Photinus pyralis with ^{32}P by the nick translation method [Molecular Cloning, pages 109-112, Cold Spring Harbor Laboratory (1982) and J. Mol. Biol., 113, 237-251 (1971)], using the colony hybridization method [Protein, Nucleic Acid & Enzyme, 26, 575-579 (1981)], an *Escherichia coli* strain having plasmid DNA containing Luciola cruciata luciferase c-DNA of 2.0 Kb can be obtained from a Luciola cruciata-derived c-DNA library prepared using plasmid pUC19 DNA (manufactured by Takara Shuzo Co., Ltd.) as a vector.

To obtain the purified plasmid DNA, there is used, for example, a method described in Proc. Natl. Acad. Sci., 62 1159-1166 (1969), etc.

By acting on the purified plasmid DNA, for example, restriction enzyme, e.g., Pst I (manufactured by Takara Shuzo Co., Ltd.) at a temperature of 30°C to 40°C, preferably at 37°C for 1 to 24 hours, preferably for 2 hours, the resulting solution obtained after completion of the reaction is subjected to agarose gel

electrophoresis [which is described in Molecular Cloning, page 150, Cold Spring Harbor Laboratory (1982)] to obtain the DNA containing the gene coding for luciferase derived from Luciola cruciata.

Next, production of the Luciola lateralis-derived luciferase-coding gene in accordance with this invention are described below.

5 Firstly, as source from which the m-RNA coding for luciferase derived from Luciola lateralis is collected, the posterior portion of Luciola lateralis is preferred since the m-RNA is present in the posterior portion of this firefly.

Preparation of m-RNA from the posterior portion of the firefly and synthesis of c-DNA from the m-RNA can be conducted, for example, in quite the same manner as in the preparation of the m-RNA of Photinus pyralis and synthesis of the c-DNA described above.

10 Then, the thus obtained c-DNA is integrated into a vector DNA, for example, plasmid pUC119 DNA, etc. to obtain various recombinant plasmid DNAs. Using these DNAs, transformation of E. coli DH1 (ATCC 33849), E. coli HB101 (ATCC 33694), etc. is effected by the method of Hanahan [DNA Cloning, 1, 109-135 (1985)] to obtain various transformants.

15 Next, after labeling c-DNA of luciferase derived from Luciola cruciata with ^{32}P by the nick translation method [Molecular Cloning, pages 109-112, Cold Spring Harbor Laboratory (1982) and J. Mol. Biol., 113, 237-251 (1977)], using the colony hybridization method [Protein, Nucleic Acid & Enzyme, 26, 575-579 (1981)], an Escherichia coli strain having plasmid DNA containing Luciola lateralis luciferase c-DNA of 2.0 Kb can be obtained from a Luciola lateralis-derived c-DNA library prepared using plasmid pUC119 DNA

20 (manufactured by Takara Shuzo Co., Ltd.) as a vector. To obtain the purified recombinant DNA from the thus obtained microorganism, there is used, for example, a method described in Proc. Natl. Acad. Sci., 62 1159-1166 (1969), etc.

By acting on the purified, new recombinant DNA, for example, 2 units of restriction enzyme Eco RI (manufactured by Takara Shuzo Co., Ltd.) at a temperature of 30 °C or higher, preferably at 37 °C, for 1 to 25 4 hours, preferably for 2 hours, partial digestion is effected. Then, digestion product is subjected to agarose gel electrophoresis to obtain 2,000 bp DNA fragment containing all the gene coding for luciferase derived from Luciola lateralis.

On the other hand, a nucleotide sequence of this luciferase gene is determined by the method as shown in item 18 in the example. The determined nucleotide sequence is shown in Fig. 6. Subsequently, an amino acid sequence of polypeptide translated from the nucleotide sequence is identified. The results are 30 shown in Fig. 7.

The gene encoding the thus identified amino acid sequence is also included in this invention.

The above-mentioned microorganism is then cultured in a medium and luciferase is collected from the culture.

35 Any medium may be used as far as it is used to culture microorganisms belonging to the genus Escherichia. Mention may be made of, for example, 1% (W/V) of trypton, 0.5% (W/V) of yeast extract, 0.5% (W/V) of NaCl and 1 mM of isopropyl- β -D-thiogalactoside, etc.

Temperature for the cultivation is between 30 and 40 °C, preferably about 37 °C and a time period for the cultivation is, for example, 4 to 8 hours, preferably about 4 hours.

40 The cells are collected from the culture by centrifugation at 8,000 r.p.m. for about 10 minutes. The obtained cells are homogenized by the method described in, for example, Methods in Enzymology, 133, 3-14 (1986) to obtain a crude enzyme solution.

The crude enzyme solution may be usable as it is; if necessary and desired, the crude enzyme solution can be purified by fractionation with ammonium sulfate, hydrophobic chromatography (for example, using BUTYL TOYOPEARL 650C, etc.), gel filtration (using, e.g., Ultrogel AcA34, etc.) thereby to give purified 45 luciferase.

Physicochemical properties of the thus obtained luciferase are as described below.

(1) Action

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The enzyme catalyzes the oxidation of luciferin by an oxygen molecule, as shown by the enzymatic reaction equation:

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Present enzyme

Luciferin + ATP + O₂



Oxyluciferin + AMP + Pyrophosphoric acid + O₂ + light

(2) Substrate specificity

The enzyme does not act on ADP, CTP, UTP and GTP.

(3) Optimum pH, and pH range for stability

The optimum pH is, as shown in Fig. 1, 7.5 to 9.5 as measured by carrying out the reaction by the use of luciferin as a substrate at various pH values of 25 mM glycylglycine buffer solution in the range of 6.5 to 11.5 and at a temperature of 30 °C, and measuring the quantity of light (the number of photons) emitted in 20 seconds. The stable pH range of the enzyme is, as shown in Fig. 2, 6.0 to 10.5 as measured by adding the enzyme to each of buffer solutions [25 mM phosphate buffer solution (pH 4.6-8.0) and 25 mM glycine.sodium chloride-sodium hydroxide buffer solutions (pH 8.0 - 11.5), each of which contains ammonium sulfate to 10% saturation] containing luciferin, and allowing the enzyme to act at a temperature of 0 °C for 4 hours. In Fig. 2, O—O and Δ—Δ show the activity in the case of using the 25 mM phosphate buffer solutions and the activity in the case of using 25 mM glycine.sodium chloride-sodium hydroxide buffer solutions, respectively.

(4) Measurement of titer

A luciferin mixed solution is prepared by mixing 8 ml of 25 mM glycylglycine buffer solution (pH 7.8), 0.5 ml of a magnesium sulfate solution [a solution prepared by adding magnesium sulfate to 25 mM glycylglycine buffer solution (pH 7.8), a magnesium sulfate concentration of 0.1 M] and 0.8 ml of a luciferin solution [a solution prepared adding luciferin to 25 mM glycylglycine buffer solution (pH 7.8), a luciferin concentration of 1 mM].

Into a mixture of 400 μl of the luciferin mixed solution thus obtained and 10 μl of luciferase to be assayed is poured 80 μl of an ATP solution [a solution prepared by adding ATP to 25 mM glycylglycine buffer solution (pH 7.8), an ATP concentration of 10 mM]. Simultaneously with the pouring, the number of photons generated is measured by adding up for 20 seconds by means of a luminometer (LUMINESCENCE READER BLR-201, manufactured by ALOKA Co., Ltd.).

(5) Range of temperature suitable for action

When the reaction is carried out at pH 7.8 at each temperature and the quantity of light (the number of photons) emitted in 20 seconds is measured, the suitable temperature for the action ranges from 0 ° to 50 °C.

(6) Conditions for inactivation by pH

At pH values of 5.0 or lower and 12.0 or higher, the enzyme is completely inactivated 4 hours after.

As is clear from the foregoing description, according to this invention, luciferase can be efficiently produced in an extremely short period of time, by culturing the microorganism belonging to the genus *Escherichia* which contains the recombinant DNA having integrated therein the *Luciola lateralis*-derived luciferase gene of this invention. Therefore, this invention is extremely useful from an industrial point of view.

Hereafter this invention will be described in more detail by referring to the examples below.

Example

In Items 1 to 10 below, production of DNA containing a gene coding for luciferase of *Photinus pyralis* as one of fireflies (this DNA is used as a probe upon survey of DNA containing a gene coding for luciferase of *Luciola cruciata*) is described. Further in Items 11 to 13 below, production of DNA containing a gene coding for luciferase derived from *Luciola cruciata* (this DNA is used as a probe upon survey of DNA containing a gene coding for luciferase of *Luciola lateralis*) is described.

1. Preparation of m-RNA

Using a mortar and a pestle, 1 g of the dry posterior portion (manufactured by Sigma Co., Ltd.) of *Photinus pyralis* as one of fireflies was thoroughly ground, to which 5 ml of dissolution buffer [20 mM Tris-hydrochloride buffer (pH 7.4)/10 mM NaCl/3 mM magnesium acetate/5% (W/V) sucrose/1.2 % (V/V) Triton X-100/10 mM vanadyl nucleoside complex (manufactured by New England Biolab Co., Ltd.)] was added. The mixture was further ground as in the manner described above to give a solution containing the ground posterior portion of *Photinus pyralis*.

In a cup blender (manufactured by Nippon Seiki Seisakusho) was charged 5 ml of the thus obtained solution. After treating at 5,000 r.p.m. for 5 minutes, 12 ml of guanidine isothiocyanate solution (6M guanidine isothiocyanate/37.5 mM sodium citrate (pH 7.0)/0.75% (W/V) sodium N-lauroylsarcosine/0.15 M β -mercaptoethanol) was added to the system. The mixture was treated with the blender described above at 3,000 r.p.m. for 10 minutes. The resulting solution was filtered using a threefold gauze to give the filtrate. The filtrate was gently poured in layers onto 4 tubes for ultracentrifuging machine (manufactured by Hitachi Koki Co., Ltd.) in which 1.2 ml each of 5.7 M cesium chloride solution had previously be laid in layers. Using the ultracentrifuging machine (manufactured by Hitachi Koki Co., Ltd., SCP55H), centrifugation was performed at 30,000 r.p.m. for 16 hours to give precipitates.

The obtained precipitates were washed with cold 70% (V/V) ethanol and suspended in 4 ml of 10 mM Tris buffer [10 mM Tris-hydrochloride (pH 7.4)/5 mM EDTA/1% sodium dodecylsulfate]. The equal amount of a mixture of n-butanol and chloroform in 1 : 4 (volume ratio) was added to the mixture to perform extraction. The extract was centrifuged at 3,000 r.p.m. for 10 minutes in a conventional manner to separate into the aqueous phase and the organic solvent phase. To the organic solvent phase was added 4 ml of 10 mM Tris buffer described above. The procedure for the extraction and separation described above was repeated twice. To the aqueous phase obtained were added a 1/10 amount of 3 M sodium acetate (pH 5.2) and a 2-fold amount of cold ethanol were added. After allowing to stand at a temperature of -20°C for 2 hours, the mixture was centrifuged at 8,000 r.p.m. for 20 minutes in a conventional manner to precipitate RNA. The obtained RNA was dissolved in 4 ml of water. After the operation for precipitation with ethanol described above was carried out, the obtained RNA was dissolved in 1 ml of water to give 3.75 mg of RNA.

By repeating the foregoing procedure again, 7 mg in total of RNA was prepared. To select m-RNA from the RNA, 7 mg of RNA was subjected to oligo(dT)-cellulose (manufactured by New England Biolab Co., Ltd.) column chromatography.

As the column, 2.5 ml of Terumo syringe (manufactured by Terumo Co., Ltd.) was used. After 0.5 g of resin was swollen with elution buffer [10 mM Tris-hydrochloride buffer (pH 7.6)/1 mM DETA/0.1% (W/V) sodium dodecylsulfate], the resin was packed in the column and equilibrated with binding buffer [10 mM Tris-hydrochloride (pH 7.6)/1 mM EDTA/0.4 M NaCl/0.1% (W/V) sodium dodecylsulfate].

To 7 mg of RNA was added the same amount of buffer [10 mM Tris-hydrochloride (pH 7.6)/1 mM EDTA/0.8 M NaCl/0.1% (W/V) sodium dodecylsulfate]. The mixture was heat-treated at a temperature of 65°C for 10 minutes and then quenched in ice water. After subjecting to oligo(dT)-cellulose column, the resin was washed with binding buffer to completely wash unbound r-RNA and t-RNA out. Further m-RNA was eluted with eluting buffer to give 40 μ g of m-RNA.

2. Concentration of luciferase m-RNA

Next, the luciferase m-RNA was concentrated by sucrose density gradient centrifugation.

Sucrose density gradient of 10 to 25% (W/V) was produced by charging 0.5 ml of 40% (W/V) sucrose solution [50 mM Tris-hydrochloride buffer (pH 7.5)/20 mM NaCl/1 mM EDTA/40% (W/V) sucrose] in a polyaroma tube for Rotor SW41 manufactured by Beckmann Co., Ltd., laying 2.4 ml each of 25% (W/V), 20% (W/V), 15% (W/V) and 10% (W/V) of the sucrose solution in layers and allowing to stand the system at a temperature of 4°C for 24 hours. To the sucrose density gradient, 30 μ g of m-RNA was laid to form a layer. Using SW41 Rotor manufactured by Beckmann Co., Ltd., centrifugation was conducted at 30,000

r.p.m. at a temperature of 18°C for 18 hours in a conventional manner. After the centrifuging operation, fractionation was performed by 0.5 ml each and m-RNA was recovered by the ethanol precipitation method. The m-RNA was dissolved in 10 µl of water.

Next, protein encoded by the m-RNA was examined, whereby the fraction concentrated on m-RNA of luciferase was identified. One microliter of the fractionated RNA, 9 µl of rabbit reticular erythrocyte lysate (manufactured by Amersham Co., Ltd.) and 1 µl of [³⁵S] methionine (manufactured by Amersham Co., Ltd.) were mixed and reacted at a temperature of 30°C for 30 minutes. To the reaction mixture was added 150 µl of NET buffer [150 mM NaCl/5 mM EDTA/0.02% (W/V) NaN₃/20 mM Tris-hydrochloride buffer (pH 7.4)-/0.05% (W/V) Nonidet P-40 (manufactured by Besesda Research Laboratories Co., Ltd., surface active agent)] and, 1 µl of antiluciferase serum (produced as will be later described) was added to the mixture. After allowing to stand at a temperature of 20°C for 30 hours, 10 mg of Protein A Sepharose (manufactured by Pharmacia Fine Chemicals Inc.) was added to the mixture. The resulting mixture was then centrifuged at 12,000 r.p.m. for a minute in a conventional manner to recover the resin.

The recovered resin was washed three times with 200 µl of NET buffer. To the resin was added 40 µl of sample buffer for SDS-PAGE [62.5 mM Tris-hydrochloride buffer (pH 6.8)/10% (V/V) glycerol/2% (W/V) sodium dodecylsulfate/5% (V/V) β-mercaptoethanol/0.02% (W/V) bromophenol blue]. The mixture was boiled at a temperature of 100°C for 3 minutes and centrifuged at 12,000 r.p.m. for a minute in a conventional manner to recover the supernatant. The whole amount was applied onto 7.5% (W/V) sodium dodecylsulfate-polyacrylamide gel.

Gel electrophoresis was performed by the method of Laemmli [Nature, 227, 680 (1970)]. After the electrophoresis, the gel was immersed in 10% (V/V) acetic acid for 30 minutes to immobilize protein. Then, the gel was immersed in water for 30 minutes and further immersed in 1 M sodium salicylate solution for 30 minutes and then dried to give a dry gel. The dry gel was subjected to fluorography using an X ray film (manufactured by Fuji Photo Film Co., Ltd.; RX).

By the foregoing procedure, the band of luciferase protein was recognized on the X ray film only in the case of using the RNA from the fraction in which the luciferase m-RNA was present and, the fraction wherein the luciferase m-RNA was concentrated could be identified.

3. Production of anti-serum

Rabbit anti-luciferase serum to purified luciferase was produced by the following method.

A luciferase solution having a 3.2 mg/ml concentration [solution obtained by dissolving luciferase manufactured by Sigma Co., Ltd. in 0.5 M glycylglycine solution (pH 7.8)], 0.7 ml, was suspended in an equal amount of Freund's complete adjuvant. 2.24 mg of the suspension was administered as an antigen to the palm of Japanese white rabbit weighing 2 kg as an antigen. After feeding for 2 weeks, the same amount of antigen as in the initial amount was intracutaneously administered to the back. After feeding for further one week, similar procedure was performed. Further one week after feeding, whole blood was collected.

The obtained blood was allowed to stand at a temperature of 4°C for 18 hours and then centrifuged at 3,000 r.p.m. for 15 minutes in a conventional manner to give anti-luciferase serum as the supernatant.

4. Synthesis of c-DNA

Synthesis of c-DNA was carried out using a kit manufactured by Amersham Co., Ltd.

Using 2 µg of m-RNA obtained as described above, synthesis of c-DNA was carried out in accordance with the methods described in Mol. Cell Biol., 2, 161 (1982) and Gene, 25, 263 (1983). As the result, 300 ng of double stranded c-DNA was obtained.

This c-DNA, 150 ng, was dissolved in 7 µl of TE buffer [10 mM Tris-hydrochloride buffer (pH 7.5)/1 mM EDTA]. To the solution were added, respectively, 11 µl of a mixture [280 mM sodium cacodylate (pH 6.8)-/60 mM Tris-hydrochloride buffer (pH 6.8)/2 mM cobalt chloride] and 3.8 µl of a tailing mixture [7.5 µl of 10 mM dithiothreitol/1 µl of 10 ng/ml poly(A)/2 µl of 5mM dCTP/110 µl of water]. Furthermore, 29 units of terminal transferase (manufactured by Boehringer Mannheim GmbH) was added to the mixture. After reacting at a temperature of 30°C for 10 minutes, 2.4 µl of 0.25 M EDTA and 2.4 µl of 10% (W/V) sodium dodecylsulfate were added to the mixture to discontinue the reaction.

The solution in which the reaction had been discontinued was subjected to a treatment for removing protein using 25 µl of water-saturated phenol. Then, 25 µl of 4 M ammonium acetate and 100 µl of cold ethanol were added to the recovered aqueous phase, respectively. The mixture was allowed to stand at a temperature of -70°C for 15 minutes and centrifuged at 12,000 r.p.m. for 10 minutes to recover c-DNA. The c-DNA was dissolved in 10 µl of TE buffer to give a c-DNA solution.

As described above, 100 ng of the c-DNA with the deoxycytidine tail was obtained.

5. Production of recombinant plasmid pMCE10 DNA used in vector

Plasmid pKN305 DNA produced by the method described in T. Masuda et al., *Agricultural Biological Chemistry*, 50, 271-279 (1986) using plasmid pBR325 (manufactured by BRL Co.) and plasmid pBR322 DNA (manufactured by Takara Shuzo Co., Ltd.), and pMC1403-3 DNA (described in Japanese Patent Publication KOKAI 61-274683) were added by 1 µg each to 10 µl of a mixture [50 mM Tris-hydrochloride buffer (pH 7.5)/10 mM MgCl₂/100 mM NaCl/1 mM dithiothreitol]. Further, 2 units each of Hind III and Sal I (both manufactured by Takara Shuzo Co., Ltd.) were added to the mixture. By reacting at a temperature of 37 °C for an hour, a cleavage treatment was effected. Extraction with phenol and precipitation with ethanol were conducted in a conventional manner to give precipitates. The precipitates were dissolved in 10 µl of ligation buffer [20 mM MgCl₂/66 mM Tris-hydrochloride buffer (pH 7.6)/1 mM ATP/15 mM dithiothreitol] to give a solution. Furthermore, 1 unit of T4 DNA ligase (manufactured by Takara Shuzo Co., Ltd.) was added thereto to perform ligation at a temperature of 20 °C for 4 hours. Then, using this reaction solution, *E. coli* JM101 (ATCC 33876) was transformed according to the transformation method described in [J. Bacteriology, 119, 1072-1074 (1974)]. By examination of chemical resistance (ampicillin resistance and tetracycline sensitivity) and β-galactosidase activity, a transformant was obtained. Recombinant plasmid DNA contained in the strain was named pMCE10. *E. coli* JM101 strain containing this recombinant plasmid DNA pMCE10 DNA was cultured in medium composed of 1% (W/V) of trypton, 0.5% (W/V) of yeast extract and 0.5% (W/V) of NaCl at a temperature of 37 °C for 16 to 24 hours. Twenty milliliters of the thus obtained culture solution of *E. coli* JM101 (pMCE10) was inoculated on 1 liter of the medium followed by shake culture at a temperature of 37 °C for 3 hours. After the addition of 0.2 g of chloramphenicol, cultivation was conducted at the same temperature for further 20 hours to give a culture solution.

Next, the culture solution was centrifuged at 6,000 r.p.m. for 10 minutes in a conventional manner to give 2 g of wet cells. After the cells were suspended in 20 ml of 350 mM Tris-hydrochloride buffer (pH 8.0) containing 25% (W/V) sucrose, 10 mg of lysozyme, 8 ml of 0.25 M EDTA solution (pH 8.0) and 8 ml of 20% (W/V) sodium dodecylsulfate were added to the suspension, respectively. The mixture was kept at a temperature of 60 °C for 30 minutes to give a lysate solution.

To the lysate solution was added 13 ml of 5 M NaCl solution. The mixture was treated at a temperature of 4 °C for 16 hours and then centrifuged at 15,000 r.p.m. in a conventional manner to give an extract. The extract was subjected to the phenol extraction and the ethanol precipitation in a conventional manner to give precipitates.

Then, the precipitates were dried under reduced pressure in a conventional manner and dissolved in 10 mM Tris-hydrochloride buffer (pH 7.5) containing 1 mM EDTA. To the solution were further added 6 g of cesium chloride and 0.2 ml of ethidium bromide solution (10 mg/ml). The resulting mixture was subjected to an equilibrated density gradient centrifugation treatment using an ultracentrifuging machine at 39,000 r.p.m. for 42 hours in a conventional manner thereby to isolate recombinant plasmid pMCE10 DNA. After ethidium bromide was removed using n-butanol, dialysis was performed to 10 mM Tris-hydrochloride buffer (pH 7.5) containing 1 mM EDTA to give 500 µg of purified recombinant plasmid pMCE10 DNA.

6. Production of vector DNA

The thus obtained recombinant plasmid pMCE10 DNA, 15 µg, was dissolved in 90 µl of TE buffer described in Item 4. After 10 µl of Med buffer [100 mM Tris-hydrochloride buffer (pH 7.5)/10 mM MgCl₂/10 mM dithiothreitol/500 mM NaCl] was added to the solution, 30 units of restriction enzyme Acc I (manufactured by Takara Shuzo Co., Ltd.) was further added to the mixture. A cleavage treatment was conducted at a temperature of 37 °C for an hour to give the cleavage product. To the cleavage product was added 100 µl of water-saturated phenol, whereby protein was removed. Then, the aqueous phase was recovered and a 1/10-fold amount of 3 M sodium acetate (pH 7.5) and a 2-fold amount of cold ethanol were added to the aqueous phase. After allowing to stand at a temperature of -70 °C for 15 minutes, the mixture was centrifuged at 12,000 r.p.m. for 10 minutes to recover DNA.

This DNA was dissolved in 10 µl of TE buffer and 15 µl of a mixture [280 mM sodium cacodylate (pH 6.8)/60 mM Tris-hydrochloride buffer (pH 6.8)/2 mM cobalt chloride] was added to the solution. Then, 5 µl of a tailing solution mixture (described in Item 4) (5 mM dGTP was used) was further added to the mixture. Furthermore, 5 units of terminal transferase (manufactured by Takara Shuzo Co., Ltd.) was added to react at a temperature of 37 °C for 15 minutes. By post-treatment in a manner similar to the c-DNA tailing reaction described in Item 4, DNA with the deoxyguanosine tail at the Acc I site of recombinant plasmid pMCE10

DNA was produced.

On the other hand, DNA with the deoxyguanosine tail at the Pst I site of plasmid pUC19 DNA was also produced at the same time.

To a solution of 30 µg of plasmid pUC19 DNA (manufactured by Takara Shuzo Co., Ltd.) in 350 µl of TE buffer were added 40 µl of Med buffer and 120 units of restriction enzyme Pst I (manufactured by Takara Shuzo Co., Ltd.). After a cleavage treatment at a temperature of 37°C for an hour, DNA was recovered by the phenol treatment for removing protein and ethanol precipitation in a conventional manner.

The obtained DNA was dissolved in 35 µl of TE buffer. To the solution were added 50 µl of a mixture [280 mM sodium cacodylate (pH 6.8)/60 mM Tris-hydrochloride buffer (pH 6.8)/1 mM cobalt chloride], 19 µl of the tailing mixture (containing dGTP instead of dCTP) described in Item 4 and 60 units of terminal transferase (manufactured by Takara Shuzo Co., Ltd.). After reacting at a temperature of 37°C for 10 minutes, DNA was recovered by the phenol treatment for removing protein and ethanol precipitation in a conventional manner.

7. Annealing and transformation

The thus synthesized c-DNA, 15 ng and 200 ng of vector DNA were dissolved in 35 µl of annealing buffer [10 mM Tris-hydrochloride buffer (pH 7.5)/100 mM NaCl/1 mM EDTA]. The solution was allowed to stand at a temperature of 65°C for 2 minutes, at a temperature of 46°C for 2 hours, at a temperature of 37°C for an hour and at a temperature of 20°C for 18 hours thereby to anneal c-DNA and vector DNA.

Using the annealed DNA, *E. coli* DH1 strain (ATCC 33849) was transformed by the method of Hanahan [DNA Cloning, 1, 109-135 (1985)] to produce a c-DNA bank containing plasmid pUC19 DNA and recombinant plasmid pMCE10 DNA as vectors, respectively.

8. Survey of luciferase c-DNA

The Acc I site of recombinant plasmid pMCE10 DNA is present at a site which codes for *E. coli* β -galactosidase gene. Therefore, c-DNA incorporated into this site forms a fused protein with β -galactosidase. Furthermore, a promoter of β -galactosidase gene of the recombinant plasmid pMCE10 DNA has been converted into a promoter of *E. coli* tryptophan gene, as described above.

96 colonies of c-DNA having recombinant plasmid pMCE10 DNA as a vector were shake cultured in 10 ml of M9 Casamino acid medium [Molecular Cloning, 440-441, Cold Spring Harbor Laboratory (1982)] supplemented with thiamine (10 µg/ml) at a temperature of 37°C for 10 hours. After collecting the cells in a conventional manner, the cells were suspended in 200 µl of sample buffer for SDS-PAGE described in Item 2. The suspension was boiled at a temperature of 100°C for 5 minutes. This suspension, 40 µl, was subjected to electrophoresis in a conventional manner using 7.5% (W/V) polyacrylamide gel. After completion of the electrophoresis, the protein developed on the gel was transferred onto a nitrocellulose filter by the western blot method [Anal. Biochem., 112, 195 (1981)]. This nitrocellulose filter was stained with anti-luciferase serum using immune blot assay kit (manufactured by Biorad Co.). The method was performed in accordance with the instruction of Biorad Co.

That is, the nitrocellulose filter was shaken in 100 ml of blocking solution [a solution obtained by dissolving 3% (W/V) gelatin in TBS buffer [20 mM Tris-hydrochloride buffer /500 mM NaCl (pH 7.5)] at a temperature of 25°C for 30 minutes. Next, this nitrocellulose filter was transferred into 25 ml of primary antibody solution [solution obtained by dissolving 1% (W/V) gelatin in TBS buffer and diluting luciferase anti-serum with the resulting solution] and shaken at a temperature of 25°C for 90 minutes, which was then transferred into 100 ml Tween-20 washing solution [solution obtained by dissolving 0.05% (W/V) Tween-20 in TBS buffer] and shaken at a temperature of 25°C for 10 minutes. This procedure was repeated twice. Then, the thus obtained nitrocellulose filter was transferred into 60 ml of secondary antibody solution [solution obtained by dissolving anti-rabbit antibody labeled with horse raddish peroxidase (manufactured by Biorad Co.) with a solution of 1% (W/V) gelatin in TBS buffer to 3000-fold (V/V)]. After shaking at a temperature of 25°C for 60 minutes, the nitrocellulose filter was washed with 100 ml of Tween-20 washing solution. The procedure described above was repeated twice. The thus obtained nitrocellulose filter was transferred into 120 ml of color forming solution [solution obtained by mixing a solution of 60 mg of 4-chloro-1-naphthol in 20 ml of cold methanol and a solution of 60 µl of 30% (V/V) hydrogen peroxide aqueous solution in 100 ml of TBS buffer] to form a color at a temperature of 25°C for 10 minutes.

As such, similar procedures were performed on 4 groups, with 96 colonies per one group. In the two groups, protein band stained with luciferase anti-serum was recognized. Next, 96 colonies belonging to the two groups were divided into 8 groups with 12 colonies each and similar procedure was repeated. A protein

that reacted with anti-luciferase serum was noted in one group. Finally, with respect to the 12 colonies contained in this group, each colony was treated in a similar manner, whereby a protein-producing colony that reacted with luciferase anti-serum was identified. By the foregoing procedure, 2 colonies containing luciferase c-DNA were obtained. From the two colonies, plasmid DNA was produced by the method described in Item 5. The obtained recombinant plasmid DNAs were named pALf2B8 and PALF3A6, respectively.

9. Survey of large luciferase c-DNA - Production of c-DNA probe

In 330 μ l of TE buffer was dissolved 100 μ g of recombinant plasmid, pALf3A6 DNA. To the solution were added 40 μ l of Low buffer [100 mM Tris-hydrochloride buffer (pH 7.5)/100 mM $MgCl_2$ /10 mM dithiothreitol], 130 units of Pst I (manufactured by Takara Shuzo Co., Ltd.) and 120 units of Sac I (manufactured by Boehringer Mannheim GmbH) to effect cleavage at a temperature of 37 °C for 1.5 hours.

The whole amount of DNA was separated by electrophoresis using 0.7% (W/V) agarose gel. The agarose gel electrophoresis was carried out in accordance with the method of T. Maniatis et al., Molecular Cloning, pages 156-161, Cold Spring Harbor Laboratory (1984). DNA band containing luciferase c-DNA was excised and put in a dialysis tube. After 2 ml of TE buffer was supplemented, the dialysis tube was sealed and DNA was eluted from the gel into the buffer by electrophoresis. An equivalent volume of water-saturated phenol was added to this solution. After agitation, the aqueous phase was recovered and DNA was recovered by precipitation with ethanol in a conventional manner.

10 μ g of the obtained DNA fragment was dissolved in TE buffer and 16 μ l of Med buffer and 64 units of Sau 3 AI (manufactured by Takara Shuzo Co., Ltd.) were added to the solution. After reacting at a temperature of 37 °C for 2 hours, the whole amount was subjected to electrophoresis using 5% (W/V) polyacrylamide gel thereby to isolate DNA fragments. The polyacrylamide gel electrophoresis was carried out in accordance with the method of A. Maxam [Methods in Enzymology, 65, 506 (1980)]. DNA fragment of 190 bp was isolated by the method as described above to give 1 μ g of Sau3 AI luciferase c-DNA fragment.

Using [α - ^{32}P] dCTP (manufactured by Amersham Co.), 1 μ g of this luciferase c-DNA was labeled according to the nick translation method. The nick translation method was performed using a kit manufactured by Takara Shuzo Co., Ltd. in accordance with the method described in J. Mol. Biol., 113, 237-251 (1977) and Molecular Cloning, pages 109-112, Cold Spring Harbor Laboratory (1982).

10. Survey of large luciferase c-DNA - Colony hybridization

Using as a probe the luciferase c-DNA fragment labelled with ^{32}P produced by the method described above, c-DNA bank of the posterior portion of *Photinus pyralis* wherein recombinant plasmid pUC19 DNA was a vector was surveyed by colony hybridization [(Protein, Nucleic Acid and Enzyme, 26, 575-579 (1981))] to give colonies having luciferase c-DNA. Recombinant plasmid DNA possessed by one of the colonies was named pALf3 and plasmid DNA was produced by the method described in Item 5. *E. coli* containing the recombinant plasmid DNA was named *E. coli* DH 1 (pALf3). The transformant has been deposited as ATCC 67462.

The recombinant plasmid pALf3 DNA described above was subjected to single digestion and double digestion using Xba I, Hind III, BamH I, Eco RI and Pst I (all manufactured by Takara Shuzo Co., Ltd.). The obtained DNA fragments were analyzed by agarose gel electrophoresis on mobility pattern. By comparing the obtained mobility pattern with standard mobility pattern of DNA fragment obtained by digesting λ phage DNA (manufactured by Takara Shuzo Co., Ltd.) with Hind III, the size of the c-DNA inserted in pALf3 was turned out to be 1,700 bp. A restriction enzyme map of the plasmid described above is shown in Fig. 3.

11. Preparation of m-RNA of *Luciola cruciata*

Ten grams of living *Luciola cruciata* (GENJI firefly, purchased from Seibu Department Store) were put in a ultra-low temperature freezer box and frozen. Each posterior portion was cut off with scissors. To 2 g of the obtained posterior portion was added 18 ml of guanidine isothiocyanate solution. According to the method described in Item 1, 1.1 mg of RNA was prepared. In accordance with the method described in Item 1, 1.1 mg of this RNA was subjected to column chromatography of oligo (dT)-cellulose to obtain 30 μ g of m-RNA for the posterior portion of *Luciola cruciata*.

12. Production of c-DNA bank of Luciola cruciata posterior portion

Synthesis of c-DNA was performed using a kit purchased from Amersham Co. in accordance with the method indicated by Amersham Co. which is described in Mol. Cell Biol., 2, 161 (1982) and Gene, 25, 263 (1983).

From 2 μ g of the Luciola cruciata posterior portion RNA, 0.9 μ g of double stranded c-DNA was synthesized. Using the method described in Item 4, a tail of polydeoxycytidine was added to 0.3 μ g of this c-DNA.

This c-DNA, 20 ng, and 500 ng of pUC19 plasmid produced in Item 6, wherein a polyguanosine tail had been added to the Pst I site thereof, were annealed in accordance with the method described in Item 7. E. coli DH 1 strain (ATCC 33849) was transformed by annealed DNA by the method of Hanahan [DNA Cloning, 1, 109-135 (1985)] thereby to produce c-DNA bank of Luciola cruciata tail.

13. Survey of luciferase c-DNA derived from Luciola cruciata

In 90 μ l of TE buffer was dissolved 10 μ g of recombinant plasmid pALf3 DNA obtained in Item 10 and, 10 μ l of Med buffer, 25 units of restriction enzyme Eco RI and 25 units of restriction enzyme Cla I (both manufactured by Takara Shuzo Co., Ltd.) were added to the solution. The reaction was performed at a temperature of 37 °C for 2 hours to cleave DNA. From the cleaved recombinant plasmid pALf3 DNA, 800 bp of Eco RI/Cla I DNA fragment containing luciferase c-DNA derived from Photinus pyralis (American firefly) was isolated in accordance with the method described in Item 9 using agarose gel electrophoresis. Thus, 1 μ g of Eco RI/Cla I DNA fragment was obtained. Using [α -³²P] dCTP (manufactured by Amersham Co.), 1 μ g of this DNA was labelled with ³²P in accordance with the nick translation method described in Item 9. Using as a probe the Eco RI/Cla I DNA fragment labeled with ³²P, c-DNA bank of the Luciola cruciata posterior portion was surveyed by the colony hybridization described in Item 10 thereby to select E. coli having luciferase c-DNA derived from Luciola cruciata. Several colonies of E. coli capable of hybridizing with the probe were obtained. Recombinant plasmid DNA possessed by one of these colonies was named pGLf1. The recombinant plasmid DNA was isolated in accordance with the method described in Item 5.

The recombinant plasmid pGLf1 DNA described above was subjected to single digestion and double digestion using Hpa I, Hind III, Eco RV, Dra I, Afl II, Hinc II, Pst I (all manufactured by Takara Shuzo Co., Ltd.) and Ssp I (manufactured by New England Biolab Co.). The obtained DNA fragments were analyzed by agarose gel electrophoresis on mobility pattern. By comparing the obtained mobility pattern with standard mobility pattern of DNA fragment obtained by digesting λ phage DNA (manufactured by Takara Shuzo Co., Ltd.) with Hind III, the size of the c-DNA inserted in pGLf1 was turned out to be 2,000 bp. A restriction enzyme map of the plasmid described above is shown in Fig. 4.

14. Preparation of m-RNA of Luciola lateralis

Five grams of living Luciola lateralis (HEIKE firefly, purchased from Kawahara Choju Trading Co., Ltd.) were put in a ultra-low temperature freezer box and frozen. Each posterior portion was cut off with scissors. To 1 g of the obtained posterior portion was added 18 ml of guanidine isothiocyanate solution. Following the method described in Item 1, 340 μ g of RNA was prepared. In accordance with the method described in Item 1, 340 μ g of the RNA was subjected to column chromatography of oligo(dT)-cellulose to obtain 6 μ g of m-RNA from the posterior portion of Luciola lateralis.

15. Production of c-DNA bank of Luciola lateralis posterior portion

Synthesis of c-DNA was performed using a kit purchased from Amersham Co.

Using 2.0 μ g of m-RNA obtained as above, c-DNA was synthesized in accordance with the method indicated by Amersham Co. which is described in Mol. Cell Biol., 2, 161 (1982) and Gene, 25, 263 (1983). As the result, 250 ng of double stranded c-DNA was obtained.

The thus obtained c-DNA, 250 ng, was subjected to methylation at the restriction enzyme Eco RI site using c-DNA cloning kit manufactured by Amersham Co. as instructed by Amersham Co. Furthermore, Eco RI linker was adhered to the both termini of the c-DNA.

After 1 μ l of Med buffer [100 mM Tris-hydrochloride buffer (pH 7.5)/100 mM MgCl₂/10 mM dithiothreitol/500 mM NaCl] was added to a solution of 100 ng of plasmid pUC119 DNA (manufactured by Takara Shuzo Co., Ltd.) in 8 μ l of water, 10 units of restriction enzyme Eco RI (manufactured by Takara Shuzo Co., Ltd.) was further added to the mixture. A cleavage treatment was conducted at a temperature of

37°C for an hour.

Subsequently, 1 µl of 1M Tris-hydrochloride buffer (pH 8.0) and 0.3 unit (1 µl) of alkaline phosphatase (manufactured by Takara Shuzo Co., Ltd.) were added to the cleavage product and the mixture was subjected to enzymatic reaction at a temperature of 65°C for an hour to effect dephosphorylation of the termini of the cleavage product. After 12 µl of water-saturated phenol was added to the dephosphorylated product to remove protein, 1 µl of 3M sodium acetate (pH 5.8) and 26 µl of cold ethanol were added to the recovered aqueous phase, respectively. The mixture was allowed to stand at a temperature of -70°C for 15 minutes. By centrifuging treatment at 12,000 r.p.m. for 5 minutes with a trace centrifuging machine (manufactured by TOMI SEIKO K.K., MRX-150) to recover DNA.

The thus obtained DNA was cleaved with restriction enzyme Eco RI and its termini were dephosphorylated. The resulting plasmid vector pUC119 DNA, 100 ng, was mixed with 250 ng of c-DNA produced in Item 15. After the mixture was suspended in 8 µl of water, 1 µl of ligation buffer [200 mM MgCl₂/660 mM Tris-hydrochloride buffer (pH 7.6)/10 mM ATP/150 mM dithiothreitol] was added to the resulting mixture. Furthermore, 1 unit of T4 DNA ligase (manufactured by Takara Shuzo Co., Ltd.) was added thereto and the mixture was allowed to stand at a temperature of 16°C for 16 hours, whereby ligation of the plasmid vector and c-DNA was performed to give the reaction product.

Using this reaction product, *E. coli* DH1 (ATCC 33849) strain was transformed by the method of Hanahan [DNA Cloning, 1, 109-135 (1985)] to produce a c-DNA bank derived from the posterior portion of *Luciola lateralis* containing plasmid pUC119 DNA as a vector.

16. Survey of luciferase c-DNA derived from *Luciola lateralis*

In 90 µl of TE buffer was dissolved 10 µg of recombinant plasmid pGLf1 DNA obtained in Item 13 and, 10 µl of Med buffer, 25 units of restriction enzyme Pst I (manufactured by Takara Shuzo Co., Ltd.) was added to the solution. The reaction was performed at a temperature of 37°C for 2 hours to cleave DNA. From the cleaved recombinant plasmid pGLf1 DNA, 2,000 bp of Pst I DNA fragment containing *Luciola cruciata*-derived luciferase c-DNA portion was isolated in accordance with the method described in Item 9 using agarose gel electrophoresis. Thus, 1 µg of Pst I DNA fragment was obtained. Using [α -³²P] dCTP (manufactured by Amersham Co.), 1 µg of this DNA was labelled with ³²P in accordance with the nick translation method described in Item 9. Using as a probe the Pst I DNA fragment labeled with ³²P, c-DNA bank of the *Luciola lateralis* posterior portion was surveyed by the colony hybridization described in Item 10 thereby to select *E. coli* bearing luciferase c-DNA derived from *Luciola lateralis*. Several colonies of *E. coli* capable of hybridizing with the probe were obtained. Plasmid DNA possessed by one of these colonies was named pHLf7. The recombinant plasmid DNA was isolated in accordance with the method described in Item 5.

The thus obtained *E. coli* DH1 (pHLf7) has been deposited in the Fermentation Research Institute, the Agency of Industrial Science and Technology, Japan, under the Budapest Treaty with the accession number FERM BP-1917.

The recombinant plasmid pHLf7 DNA described above was subjected to single digestion and double digestion using Hpa I, Eco RV, Apa I, Hind III and Eco RI (all manufactured by Takara Shuzo Co., Ltd.) and Ssp I (manufactured by New England Biolab Co.). The obtained DNA fragments were analyzed by agarose gel electrophoresis on mobility pattern. By comparing the obtained mobility pattern with standard mobility pattern of DNA fragment obtained by digesting λ phage DNA (manufactured by Takara Shuzo Co., Ltd.) with Hind III, the size of the gene encoding *Luciola lateralis*-derived luciferase was turned out to be 2,000 bp. A restriction enzyme map of the plasmid described above is shown in Fig. 5.

To a solution of 10 µg of recombinant plasmid pHLf7 DNA in 45 µl of TE buffer were added 5 µl of Med buffer and 2 units of restriction enzyme Eco RI (manufactured by Takara Shuzo Co., Ltd.), respectively. The mixture was reacted at a temperature of 37°C for 2 hours to give a partial digestion product of DNA.

Then, the partial digestion product was subjected to agarose gel electrophoresis described in Item 9 and 1 µg of Eco RI fragment of 2,000 bp containing all the gene encoding *Luciola lateralis*-derived luciferase was isolated.

17. Cultivation of *E. coli* DH1 (pHLf7) (FERM BP-1917) and production of crude enzyme solution

E. coli DH1 (pHLf7) (FERM BP-1917) was shake cultured in 3 ml of LB-amp medium [1% (W/V) bactotrypton, 0.5% (W/V) yeast extract, 0.5% (W/V) NaCl and ampicillin (50 µg/ml)] at a temperature of 37°C for 18 hours. This culture solution, 0.5 ml, was inoculated on 10 µl of the aforesaid LB-amp medium

and 1 mM isopropyl- β -D-thiogalactoside was added thereto. After shake culture at a temperature of 37°C for 4 hours, the culture was subjected to a centrifuging operation at 8,000 r.p.m. for 10 minutes to give 20 mg of wet cells.

The recovered cells were suspended in 0.9 ml of a buffer composed of 0.1 M KH_2PO_4 (pH 7.8), 2 mM EDTA, 1 mM dithiothreitol and 0.2 mg/ml protamine sulfate. Further 100 μl of 10 mg/ml lysozyme solution was supplemented to the suspension. The mixture was allowed to stand in ice for 15 minutes. Next, the suspension was frozen in methanol-dry ice bath and then allowed to stand at a temperature of 25°C to completely thaw. Further by performing a centrifuging operation at 12,000 r.p.m. for 5 minutes, 1 ml of crude enzyme solution was obtained as the supernatant.

The luciferase activity in the thus obtained crude enzyme solution was performed by the method described below. The results are shown in Table 1 below.

The measurement of luciferase activity in the crude enzyme solution obtained was performed by counting the number of photons generated in accordance with the method of Kricka [Archives of Biochemistry and Biophysics, 217, 674 (1982)].

That is, 260 μl of 25 mM glycylglycine buffer (pH 7.8), 16 μl of 0.1 M magnesium sulfate and 24 μl of 1 mM luciferine (manufactured by Sigma Inc.) and 10 μl of the crude enzyme solution were mixed. Then 100 μl of 20 mM ATP was added to the mixture. The number of photons generated was integrated for 20 seconds. The integrated values are shown in Table 1 below. For purpose of comparison, a luciferase activity was measured also with plasmid pUC119 DNA-bearing *E. coli* DH1 strain [*E. coli* DH1 (pUC119)]. The results are also shown in Table 1 below.

Table 1

25	Item	
	<u>Sample</u>	<u>Number of Photon/ml Culture Solution</u>
30	<u>E. coli</u> DH1 (pHLf7) (invention)	6.2×10^6
35	<u>E. coli</u> DH1 (pUC119) (control)	1.0×10^4

As is clear from the table above, it is noted that the count of photons increased in *E. coli* DH1 (pHLf7) bearing the recombinant plasmid pHLf7 containing the luciferase gene of this invention as compared to the control and therefore, luciferase is produced in the cells of *E. coli* used in this invention.

18. Analysis of nucleotide sequence of luciferase c-DNA derived from *Luciola lateralis*

Recombinant plasmid pHLf7 DNA, 10 μg , was cleaved with restriction enzyme Eco RI (manufactured by Takara Shuzo Co., Ltd.) to give 2.0 μg of 1.7 Kb DNA fragment and 0.5 μg of 0.3 Kb DNA fragment, containing luciferase c-DNA. These DNA fragments were subcloned at the Eco RI site of plasmid pUC118 DNA (manufactured by Takara Shuzo Co., Ltd.) to give 4 plasmids, pHLf11, pHLf12, pHLf13 and pHLf14, based on differences in kind of the inserted fragments (1.7 Kb and 0.3 Kb) and in orientation of the insertion (the 1.7 Kb fragment was subcloned to pHLf11 and pHLf12, and the 0.3 Kb fragment was subcloned to pHLf13 and pHLf14).

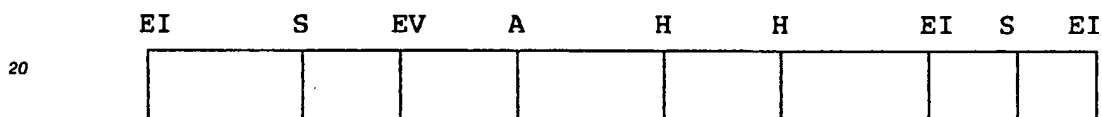
Cleavage treatment of the recombinant plasmids pHLf7 DNA and plasmid pUC118 DNA with Eco RI (method described in Item 6), isolation of the luciferase c-DNA fragment using agarose gel electrophoresis (method described in Item 9), ligation of plasmid pUC119 DNA and the luciferase c-DNA fragment (method described in Item 5), transformation of *E. coli* JM101 strain (ATCC 33876) using ligation reaction liquid (method described in Item 5) and production of recombinant plasmids pHLf11, pHLf12, pHLf13 and pHLf14 (method described in Item 5) followed the methods described within parentheses.

Next, using the recombinant plasmid DNAs pHLf11, pHLf12, pHLf13 and pHLf14, plasmid DNAs wherein various deletions were introduced into the luciferase c-DNA were produced using a deletion kit for

kilobase sequence (manufactured by Takara Shuzo Co., Ltd.) in accordance with the method of Henikoff [Gene, 28, 351-359 (1984)]. These plasmid DNAs were introduced into *E. coli* JM101 strain (ATCC 33876) by the method described in Item 5. By infecting the thus obtained *E. coli* with helper phage M13K07 (manufactured by Takara Shuzo Co., Ltd.), single strand DNA was produced in accordance with the method of Messing [Methods in Enzymology, 101, 20-78 (1983)]. Sequencing with the obtained single strand DNA was carried out by the method of Messing described above, using M13 sequencing kit (manufactured by Takara Shuzo Co., Ltd.). Gel electrophoresis for analyzing a nucleotide sequence was carried out using 8% (W/V) polyacrylamide gel (manufactured by Fuji Photo Film Co., Ltd.). The nucleotide sequence of the *Luciola lateralis*-derived luciferase-coding c-DNA obtained is shown in Fig. 6. Fig. 7 shows an amino acid sequence of polypeptide translated from the c-DNA and Fig. 8 shows a sequence corresponding to c-DNA in the amino acid sequence.

Claims

1. A *Luciola lateralis*-derived luciferase gene defined by a restriction enzyme map described below:



wherein EI represents Eco RI, S represents Ssp I, EV represents Eco RV, A represents Apa I and H represents Hpa I.

2. A luciferase gene according to claim 1, which encodes an amino acid sequence shown below:

5	Met Glu Asn Met Glu Asn Asp Glu Asn Ile	10
	Val Tyr Gly Pro Glu Pro Phe Tyr Pro Ile	20
	Glu Glu Gly Ser Ala Gly Ala Gln Leu Arg	30
10	Lys Tyr Met Asp Arg Tyr Ala Lys Leu Gly	40
	Ala Ile Ala Phe Thr Asn Ala Leu Thr Gly	50
15	Val Asp Tyr Thr Tyr Ala Glu Tyr Leu Glu	60
	Lys Ser Cys Cys Leu Gly Glu Ala Leu Lys	70
20	Asn Tyr Gly Leu Val Val Asp Gly Arg Ile	80
	Ala Leu Cys Ser Glu Asn Cys Glu Glu Phe	90
25	Phe Ile Pro Val Leu Ala Gly Leu Phe Ile	100
	Gly Val Gly Val Ala Pro Thr Asn Glu Ile	110
30	Tyr Thr Leu Arg Glu Leu Val His Ser Leu	120
	Gly Ile Ser Lys Pro Thr Ile Val Phe Ser	130
35	Ser Lys Lys Gly Leu Asp Lys Val Ile Thr	140
	Val Gln Lys Thr Val Thr Ala Ile Lys Thr	150
40	Ile Val Ile Leu Asp Ser Lys Val Asp Tyr	160
	Arg Gly Tyr Gln Ser Met Asp Asn Phe Ile	170
45	Lys Lys Asn Thr Pro Gln Gly Phe Lys Gly	180
	Ser Ser Phe Lys Thr Val Glu Val Asn Arg	190
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	Lys	Glu	Gln	Val	Ala	Leu	Ile	Met	Asn	Ser	200
											210
5	Ser	Gly	Ser	Thr	Gly	Leu	Pro	Lys	Gly	Val	
											220
	Gln	Leu	Thr	His	Glu	Asn	Ala	Val	Thr	Arg	
											230
10	Phe	Ser	His	Ala	Arg	Asp	Pro	Ile	Tyr	Gly	
											240
	Asn	Gln	Val	Ser	Pro	Gly	Thr	Ala	Ile	Leu	
											250
15	Thr	Val	Val	Pro	Phe	His	His	Gly	Phe	Gly	
											260
	Met	Phe	Thr	Thr	Leu	Gly	Tyr	Leu	Thr	Cys	
											270
20	Gly	Phe	Arg	Ile	Val	Met	Leu	Thr	Lys	Phe	
											280
	Asp	Glu	Glu	Thr	Phe	Leu	Lys	Thr	Leu	Gln	
											290
25	Asp	Tyr	Lys	Cys	Ser	Ser	Val	Ile	Leu	Val	
											300
	Pro	Thr	Leu	Phe	Ala	Ile	Leu	Asn	Arg	Ser	
											310
30	Glu	Leu	Leu	Asp	Lys	Tyr	Asp	Leu	Ser	Asn	
											320
	Leu	Val	Glu	Ile	Ala	Ser	Gly	Gly	Ala	Pro	
											330
	Leu	Ser	Lys	Glu	Ile	Gly	Glu	Ala	Val	Ala	
35											340
	Arg	Arg	Phe	Asn	Leu	Pro	Gly	Val	Arg	Gln	
											350
	Gly	Tyr	Gly	Leu	Thr	Glu	Thr	Thr	Ser	Ala	
40											360
	Ile	Ile	Ile	Thr	Pro	Glu	Gly	Asp	Asp	Lys	
											370
	Pro	Gly	Ala	Ser	Gly	Lys	Val	Val	Pro	Leu	
45											380
	Phe	Lys	Ala	Lys	Val	Ile	Asp	Leu	Asp	Thr	

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	Lys	Lys	Thr	Leu	Gly	Pro	Asn	Arg	Arg	Gly	390
	Glu	Val	Cys	Val	Lys	Gly	Pro	Met	Leu	Met	400
5	Lys	Gly	Tyr	Val	Asp	Asn	Pro	Glu	Ala	Thr	410
	Arg	Glu	Ile	Ile	Asp	Glu	Glu	Gly	Trp	Leu	420
10	His	Thr	Gly	Asp	Ile	Gly	Tyr	Tyr	Asp	Glu	430
	Glu	Lys	His	Phe	Phe	Ile	Val	Asp	Arg	Leu	440
15	Lys	Ser	Leu	Ile	Lys	Tyr	Lys	Gly	Tyr	Gln	450
	Val	Pro	Pro	Ala	Glu	Leu	Glu	Ser	Val	Leu	460
20	Leu	Gln	His	Pro	Asn	Ile	Phe	Asp	Ala	Gly	470
	Val	Ala	Gly	Val	Pro	Asp	Pro	Ile	Ala	Gly	480
25	Glu	Leu	Pro	Gly	Ala	Val	Val	Val	Leu	Glu	490
	Lys	Gly	Lys	Ser	Met	Thr	Glu	Lys	Glu	Val	500
30	Met	Asp	Tyr	Val	Ala	Ser	Gln	Val	Ser	Asn	510
	Ala	Lys	Arg	Leu	Arg	Gly	Gly	Val	Arg	Phe	520
35	Val	Asp	Glu	Val	Pro	Lys	Gly	Leu	Thr	Gly	530
	Lys	Ile	Asp	Gly	Lys	Ala	Ile	Arg	Glu	Ile	540
40	Leu	Lys	Lys	Pro	Val	Ala	Lys	Met			

45 3. A luciferase gene according to Claim 1 or 2 which is represented by a nucleotide sequence shown below.

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	ATG GAA AAC ATG GAG AAC GAT GAA AAT ATT	30
5	GTG TAT GGT CCT GAA CCA TTT TAC CCT ATT	60
	GAA GAG GGA TCT GCT GGA GCA CAA TTG CGC	90
10	AAG TAT ATG GAT CGA TAT GCA AAA CTT GGA	120
	GCA ATT GCT TTT ACT AAC GCA CTT ACC GGT	150
15	GTC GAT TAT ACG TAC GCC GAA TAC TTA GAA	180
	AAA TCA TGC TGT CTA GGA GAG GCT TTA AAG	210
20	AAT TAT GGT TTG GTT GTT GAT GGA AGA ATT	240
	GCG TTA TGC AGT GAA AAC TGT GAA GAA TTC	270
25	TTT ATT CCT GTA TTA GCC GGT TTA TTT ATA	300
	GGT GTC GGT GTG GCT CCA ACT AAT GAG ATT	330
30	TAC ACT CTA CGT GAA TTG GTT CAC AGT TTA	360
	GGC ATC TCT AAG CCA ACA ATT GTA TTT AGT	390
35	TCT AAA AAA GGA TTA GAT AAA GTT ATA ACT	420
	GTA CAA AAA ACG GTA ACT GCT ATT AAA ACC	450
40	ATT GTT ATA TTG GAC AGC AAA GTG GAT TAT	480
	AGA GGT TAT CAA TCC ATG GAC AAC TTT ATT	510
45	AAA AAA AAC ACT CCA CAA GGT TTC AAA GGA	540
	TCA AGT TTT AAA ACT GTA GAA GTT AAC CGC	570

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	AAA GAA CAA GTT GCT CTT ATA ATG AAC TCT	600
5	TCG GGT TCA ACC GGT TTG CCA AAA GGT GTG	630
	CAA CTT ACT CAT GAA AAT GCA GTC ACT AGA	660
10	TTT TCT CAC GCT AGA GAT CCA ATT TAT GGA	690
	AAC CAA GTT TCA CCA GGC ACG GCT ATT TTA	720
15	ACT GTA GTA CCA TTC CAT CAT GGT TTT GGT	750
	ATG TTT ACT ACT TTA GGC TAT CTA ACT TGT	780
20	GGT TTT CGT ATT GTC ATG TTA ACG AAA TTT	810
	GAC GAA GAG ACT TTT TTA AAA ACA CTG CAA	840
25	GAT TAC AAA TGT TCA AGC GTT ATT CTT GTA	870
	CCG ACT TTG TTT GCA ATT CTT AAT AGA AGT	900
30	GAA TTA CTC GAT AAA TAT GAT TTA TCA AAT	930
	TTA GTT GAA ATT GCA TCT GGC GGA GCA CCT	960
35	TTA TCT AAA GAA ATT GGT GAA GCT GTT GCT	990
	AGA CGT TTT AAT TTA CCG GGT GTT CGT CAA	1020
40	GGC TAT GGT TTA ACA GAA ACA ACC TCT GCA	1050
	ATT ATT ATC ACA CCG GAA GGC GAT GAT AAA	1080
45	CCA GGT GCT TCT GGC AAA GTT GTG CCA TTA	1110
	TTT AAA GCA AAA GTT ATC GAT CTT GAT ACT	1140

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      AAA AAA ACT TTG GGC CCG AAC AGA CGT GGA      1 1 7 0
      GAA GTT TGT GTA AAG GGT CCT ATG CTT ATG      1 2 0 0
5      AAA GGT TAT GTA GAT AAT CCA GAA GCA ACA      1 2 3 0
      AGA GAA ATC ATA GAT GAA GAA GGT TGG TTG      1 2 6 0
10     CAC ACA GGA GAT ATT GGG TAT TAC GAT GAA      1 2 9 0
      GAA AAA CAT TTC TTT ATC GTG GAT CGT TTG      1 3 2 0
15     AAG TCT TTA ATC AAA TAC AAA GGA TAT CAA      1 3 5 0
      GTA CCA CCT GCT GAA TTA GAA TCT GTT CTT      1 3 8 0
20     TTG CAA CAT CCA AAT ATT TTT GAT GCC GGC      1 4 1 0
      GTT GCT GGC GTT CCA GAT CCT ATA GCT GGT      1 4 4 0
25     GAG CTT CCG GGA GCT GTT GTT GTA CTT GAA      1 4 7 0
      AAA GGA AAA TCT ATG ACT GAA AAA GAA GTA      1 5 0 0
30     ATG GAT TAC GTT GCT AGT CAA GTT TCA AAT      1 5 3 0
      GCA AAA CGT TTG CGT GGT GGT GTC CGT TTT      1 5 6 0
35     GTG GAC GAA GTA CCT AAA GGT CTC ACT GGT      1 5 9 0
      AAA ATT GAC GGT AAA GCA ATT AGA GAA ATA      1 6 2 0
40     CTG AAG AAA CCA GTT GCT AAG ATG

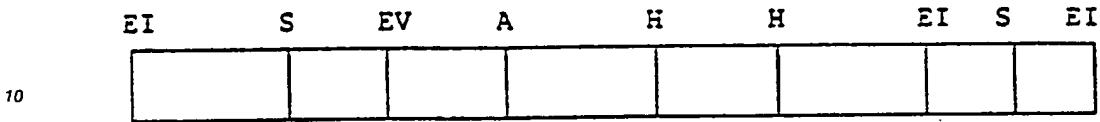
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- 45 4. A novel recombinant DNA comprising a vector DNA into which a gene coding for Luciola lateralis-derived luciferase according to claims 1-3 is inserted.
5. A novel recombinant DNA according to claim 4, wherein said vector DNA is plasmid pUC119.
- 50 6. A method for producing a luciferase which comprises culturing in a medium a microorganism belonging to the genus Escherichia and bearing a recombinant DNA according to claim 4 obtained by inserting a gene coding for Luciola lateralis-derived luciferase into a vector DNA and collecting said luciferase from the culture.
- 55 7. A method for producing a luciferase according to claim 6, wherein said vector DNA is plasmid pUC119.

Patentansprüche

1. Ein von *Luciola lateralis* abgeleitetes Luciferase-Gen, definiert durch die nachfolgend angegebene Restriktionsenzymkarte:

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worin EI Eco RI bedeutet, S Ssp I bedeutet, EV Eco RV bedeutet, A Apa I bedeutet und H Hpa I bedeutet.

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2. Luciferase-Gen nach Anspruch 1, dadurch gekennzeichnet, daß es die nachfolgend angegebene Aminosäuresequenz codiert:

20

Met Glu Asn Met Glu Asn Asp Glu Asn Ile 10

Val Tyr Gly Pro Glu Pro Phe Tyr Pro Ile 20

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Glu Glu Gly Ser Ala Gly Ala Gln Leu Arg 30

Lys Tyr Met Asp Arg Tyr Ala Lys Leu Gly 40

30

Ala Ile Ala Phe Thr Asn Ala Leu Thr Gly 50

Val Asp Tyr Thr Tyr Ala Glu Tyr Leu Glu 60

35

Lys Ser Cys Cys Leu Gly Glu Ala Leu Lys 70

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	Asn	Tyr	Gly	Leu	Val	Val	Asp	Gly	Arg	Ile	80
5	Ala	Leu	Cys	Ser	Glu	Asn	Cys	Glu	Glu	Phe	90
	Phe	Ile	Pro	Val	Leu	Ala	Gly	Leu	Phe	Ile	100
10	Gly	Val	Gly	Val	Ala	Pro	Thr	Asn	Glu	Ile	110
	Tyr	Thr	Leu	Arg	Glu	Leu	Val	His	Ser	Leu	120
15	Gly	Ile	Ser	Lys	Pro	Thr	Ile	Val	Phe	Ser	130
	Ser	Lys	Lys	Gly	Leu	Asp	Lys	Val	Ile	Thr	140
20	Val	Gln	Lys	Thr	Val	Thr	Ala	Ile	Lys	Thr	150
	Ile	Val	Ile	Leu	Asp	Ser	Lys	Val	Asp	Tyr	160
25	Arg	Gly	Tyr	Gln	Ser	Met	Asp	Asn	Phe	Ile	170
	Lys	Lys	Asn	Thr	Pro	Gln	Gly	Phe	Lys	Gly	180
30	Ser	Ser	Phe	Lys	Thr	Val	Glu	Val	Asn	Arg	190
	Lys	Glu	Gln	Val	Ala	Leu	Ile	Met	Asn	Ser	200
35	Ser	Gly	Ser	Thr	Gly	Leu	Pro	Lys	Gly	Val	210
	Gln	Leu	Thr	His	Glu	Asn	Ala	Val	Thr	Arg	220
40	Phe	Ser	His	Ala	Arg	Asp	Pro	Ile	Tyr	Gly	230
	Asn	Gln	Val	Ser	Pro	Gly	Thr	Ala	Ile	Leu	240
45	Thr	Val	Val	Pro	Phe	His	His	Gly	Phe	Gly	250
	Met	Phe	Thr	Thr	Leu	Gly	Tyr	Leu	Thr	Cys	260
50	Gly	Phe	Arg	Ile	Val	Met	Leu	Thr	Lys	Phe	270
	Asp	Glu	Glu	Thr	Phe	Leu	Lys	Thr	Leu	Gln	280

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	Asp Tyr Lys Cys Ser Ser Val Ile Leu Val	290
	Pro Thr Leu Phe Ala Ile Leu Asn Arg Ser	300
5	Glu Leu Leu Asp Lys Tyr Asp Leu Ser Asn	310
	Leu Val Glu Ile Ala Ser Gly Gly Ala Pro	320
10	Leu Ser Lys Glu Ile Gly Glu Ala Val Ala	330
	Arg Arg Phe Asn Leu Pro Gly Val Arg Gln	340
15	Gly Tyr Gly Leu Thr Glu Thr Thr Ser Ala	350
	Ile Ile Ile Thr Pro Glu Gly Asp Asp Lys	360
20	Pro Gly Ala Ser Gly Lys Val Val Pro Leu	370
	Phe Lys Ala Lys Val Ile Asp Leu Asp Thr	380
25	Lys Lys Thr Leu Gly Pro Asn Arg Arg Gly	390
	Glu Val Cys Val Lys Gly Pro Met Leu Met	400
30	Lys Gly Tyr Val Asp Asn Pro Glu Ala Thr	410
	Arg Glu Ile Ile Asp Glu Glu Gly Trp Leu	420
35	His Thr Gly Asp Ile Gly Tyr Tyr Asp Glu	430
	Glu Lys His Phe Phe Ile Val Asp Arg Leu	440
40	Lys Ser Leu Ile Lys Tyr Lys Gly Tyr Gln	450
	Val Pro Pro Ala Glu Leu Glu Ser Val Leu	460
45	Leu Gln His Pro Asn Ile Phe Asp Ala Gly	470
	Val Ala Gly Val Pro Asp Pro Ile Ala Gly	480
50	Glu Leu Pro Gly Ala Val Val Val Leu Glu	490
	Lys Gly Lys Ser Met Thr Glu Lys Glu Val	500

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Met Asp Tyr Val Ala Ser Gln Val Ser Asn 310
 5 Ala Lys Arg Leu Arg Gly Gly Val Arg Phe 320
 Val Asp Glu Val Pro Lys Gly Leu Thr Gly 330
 10 Lys Ile Asp Gly Lys Ala Ile Arg Glu Ile 340
 Leu Lys Lys Pro Val Ala Lys Met

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3. Luciferase-Gen gemäß Anspruch 1 oder 2, dadurch gekennzeichnet, daß es durch die nachfolgend angegebene Nucleotidsequenz dargestellt wird.

20 ATG GAA AAC ATG GAG AAC GAT GAA AAT ATT 350
 GTG TAT GGT CCT GAA CCA TTT TAC CCT ATT 360
 25 GAA GAG GGA TCT GCT GGA GCA CAA TTG CGC 370
 AAG TAT ATG GAT CGA TAT GCA AAA CTT GGA 380
 30 GCA ATT. GCT TTT ACT AAC GCA CTT ACC GGT 390
 GTC GAT TAT ACG TAC GCC GAA TAC TTA GAA 400
 35 AAA TCA TGC TGT CTA GGA GAG GCT TTA AAG 410
 AAT TAT GGT TTG GTT GTT GAT GGA AGA ATT 420
 40 GCG TTA TGC AGT GAA AAC TGT GAA GAA TTC 430
 TTT ATT CCT GTA TTA GCC GGT TTA TTT ATA 440
 45 GGT GTC GGT GTG GCT CCA ACT AAT GAG ATT 450

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	TAC	ACT	CTA	CGT	GAA	TTG	GTT	CAC	AGT	TTA	360
5	GGC	ATC	TCT	AAG	CCA	ACA	ATT	GTA	TTT	AGT	390
	TCT	AAA	AAA	GGA	TTA	GAT	AAA	GTT	ATA	ACT	420
10	GTA	CAA	AAA	ACG	GTA	ACT	GCT	ATT	AAA	ACC	450
	ATT	GTT	ATA	TTG	GAC	AGC	AAA	GTG	GAT	TAT	480
15	AGA	GGT	TAT	CAA	TCC	ATG	GAC	AAC	TTT	ATT	510
	AAA	AAA	AAC	ACT	CCA	CAA	GGT	TTC	AAA	GGA	540
20	TCA	AGT	TTT	AAA	ACT	GTA	GAA	GTT	AAC	CGC	570
	AAA	GAA	CAA	GTT	GCT	CTT	ATA	ATG	AAC	TCT	600
25	TCG	GGT	TCA	ACC	GGT	TTG	CCA	AAA	GGT	GTG	630
	CAA	CTT	ACT	CAT	GAA	AAT	GCA	GTC	ACT	AGA	660
30	TTT	TCT	CAC	GCT	AGA	GAT	CCA	ATT	TAT	GGA	690
	AAC	CAA	GTT	TCA	CCA	GGC	ACG	GCT	ATT	TTA	720
35	ACT	GTA	GTA	CCA	TTC	CAT	CAT	GGT	TTT	GGT	750
	ATG	TTT	ACT	ACT	TTA	GGC	TAT	CTA	ACT	TGT	780
40	GGT	TTT	CGT	ATT	GTC	ATG	TTA	ACG	AAA	TTT	810
	GAC	GAA	GAG	ACT	TTT	TTA	AAA	ACA	CTG	CAA	840
45	GAT	TAC	AAA	TGT	TCA	AGC	GTT	ATT	CTT	GTA	870
	CCG	ACT	TTG	TTT	GCA	ATT	CTT	AAT	AGA	AGT	900
50	GAA	TTA	CTC	GAT	AAA	TAT	GAT	TTA	TCA	AAT	930
	TTA	GTT	GAA	ATT	GCA	TCT	GGC	GGA	GCA	CCT	960

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5 TTA TCT AAA GAA ATT GGT GAA GCT GTT GCT 190
 AGA CGT TTT AAT TTA CCG GGT GTT CGT CAA 1020
 GGC TAT GGT TTA ACA GAA ACA ACC TCT GCA 1050
 10 ATT ATT ATC ACA CCG GAA GGC GAT GAT AAA 1080
 CCA GGT GCT TCT GGC AAA GTT GTG CCA TTA 1110
 15 TTT AAA GCA AAA GTT ATC GAT CTT GAT ACT 1140
 AAA AAA ACT TTG GGC CCG AAC AGA CGT GGA 1170
 20 GAA GTT TGT GTA AAG GGT CCT ATG CTT ATG 1200
 AAA GGT TAT GTA GAT AAT CCA GAA GCA ACA 1230
 25 AGA GAA ATC ATA GAT GAA GAA GGT TGG TTG 1260
 CAC ACA GGA GAT ATT GGG TAT TAC GAT GAA 1290
 30 GAA AAA CAT TTC TTT ATC GTG GAT CGT TTG 1320
 AAG TCT TTA ATC AAA TAC AAA GGA TAT CAA 1350
 35 GTA CCA CCT GCT GAA TTA GAA TCT GTT CTT 1380
 TTG CAA CAT CCA AAT ATT TTT GAT GCC GGC 1410
 GTT GCT GGC GTT CCA GAT CCT ATA GCT GGT 1440
 40 GAG CTT CCG GGA GCT GTT GTT GTA CTT GAA 1470
 AAA GGA AAA TCT ATG ACT GAA AAA GAA GTA 1500
 45 ATG GAT TAC GTT GCT AGT CAA GTT TCA AAT 1530
 GCA AAA CGT TTG CGT GGT GGT GTC CGT TTT 1560

50

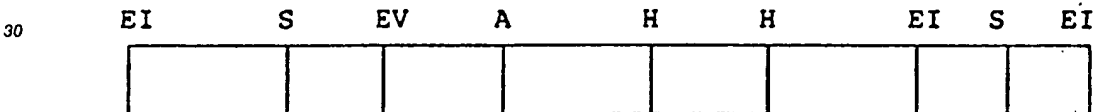
55

5 GTG GAC GAA GTA CCT AAA GGT CTC ACT GGT
 AAA ATT GAC GGT AAA GCA ATT AGA GAA ATA
 CTG AAG AAA CCA GTT GCT AAG ATG

- 10
4. Neue rekombinante DNA, umfassend eine Vektor-DNA, in die ein Gen gemäß den Ansprüchen 1 bis 3, das für von *Luciola lateralis* abgeleitete Luciferase codiert, eingeführt ist.
- 15
5. Neue rekombinante DNA gemäß Anspruch 4, dadurch gekennzeichnet, daß die Vektor-DNA Plasmid pUC119 ist.
- 20
6. Verfahren zur Herstellung einer Luciferase, dadurch gekennzeichnet, daß man in einem Medium einen Mikroorganismus der Gattung *Escherichia*, der eine rekombinante DNA gemäß Anspruch 4 trägt, die erhalten wurde durch Einführung eines Gens, das für von *Luciola lateralis* abgeleitete Luciferase codiert, in eine Vektor-DNA kultiviert und die Luciferase aus der Kultur gewinnt.
7. Verfahren zur Herstellung einer Luciferase gemäß Anspruch 6, dadurch gekennzeichnet, daß die Vektor-DNA Plasmid pUC119 ist.

25 **Revendications**

1. Gène de luciférase dérivé de *Luciola lateralis*, défini par la carte de restriction décrite ci-dessous:



35 dans laquelle EI représente EcoRI, S représente Ssp I, EV représente Eco RV, A représente Apa I et H représente Hpa I.

- 40
2. Gène de luciférase selon la revendication 1, qui code la séquence d'acides aminés présentée ci-dessous:
- 45
- 50
- 55

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10
Met Glu Asn Met Glu Asn Asp Glu Asn Ile
5 20
Val Tyr Gly Pro Glu Pro Phe Tyr Pro Ile
30
10 Glu Glu Gly Ser Ala Gly Ala Gln Leu Arg
40
Lys Tyr Met Asp Arg Tyr Ala Lys Leu Gly
50
15 Ala Ile Ala Phe Thr Asn Ala Leu Thr Gly
60
Val Asp Tyr Thr Tyr Ala Glu Tyr Leu Glu
20 70
Lys Ser Cys Cys Leu Gly Glu Ala Leu Lys
80
25 Asn Tyr Gly Leu Val Val Asp Gly Arg Ile
90
Ala Leu Cys Ser Glu Asn Cys Glu Glu Phe

30

35

40

45

50

55

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		100
	Phe Ile Pro Val Leu Ala Gly Leu Phe Ile	
5		110
	Gly Val Gly Val Ala Pro Thr Asn Glu Ile	
		120
10	Tyr Thr Leu Arg Glu Leu Val His Ser Leu	
		130
	Gly Ile Ser Lys Pro Thr Ile Val Phe Ser	
		140
15	Ser Lys Lys Gly Leu Asp Lys Val Ile Thr	
		150
	Val Gln Lys Thr Val Thr Ala Ile Lys Thr	
20		160
	Ile Val Ile Leu Asp Ser Lys Val Asp Tyr	
		170
25	Arg Gly Tyr Gln Ser Met Asp Asn Phe Ile	
		180
	Lys Lys Asn Thr Pro Gln Gly Phe Lys Gly	
		190
30	Ser Ser Phe Lys Thr Val Glu Val Asn Arg	
		200
	Lys Glu Gln Val Ala Leu Ile Met Asn Ser	
35		210
	Ser Gly Ser Thr Gly Leu Pro Lys Gly Val	
		220
40	Gln Leu Thr His Glu Asn Ala Val Thr Arg	
		230
	Phe Ser His Ala Arg Asp Pro Ile Tyr Gly	
		240
45	Asn Gln Val Ser Pro Gly Thr Ala Ile Leu	
		250
	Thr Val Val Pro Phe His His Gly Phe Gly	
50		260
	Met Phe Thr Thr Leu Gly Tyr Leu Thr Cys	

55

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		270
	Gly Phe Arg Ile Val Met Leu Thr Lys Phe	
5		280
	Asp Glu Glu Thr Phe Leu Lys Thr Leu Gln	
		290
10	Asp Tyr Lys Cys Ser Ser Val Ile Leu Val	
		300
	Pro Thr Leu Phe Ala Ile Leu Asn Arg Ser	
		310
15	Glu Leu Leu Asp Lys Tyr Asp Leu Ser Asn	
		320
	Leu Val Glu Ile Ala Ser Gly Gly Ala Pro	
20		330
	Leu Ser Lys Glu Ile Gly Glu Ala Val Ala	
		340
25	Arg Arg Phe Asn Leu Pro Gly Val Arg Gln	
		350
	Gly Tyr Gly Leu Thr Glu Thr Thr Ser Ala	
		360
30	Ile Ile Ile Thr Pro Glu Gly Asp Asp Lys	
		370
	Pro Gly Ala Ser Gly Lys Val Val Pro Leu	
35		380
	Phe Lys Ala Lys Val Ile Asp Leu Asp Thr	
		390
40	Lys Lys Thr Leu Gly Pro Asn Arg Arg Gly	
		400
	Glu Val Cys Val Lys Gly Pro Met Leu Met	
		410
45	Lys Gly Tyr Val Asp Asn Pro Glu Ala Thr	
		420
	Arg Glu Ile Ile Asp Glu Glu Gly Trp Leu	
50		430
	His Thr Gly Asp Ile Gly Tyr Tyr Asp Glu	

55

440
 Glu Lys His Phe Phe Ile Val Asp Arg Leu
 450
 5 Lys Ser Leu Ile Lys Tyr Lys Gly Tyr Gln
 460
 Val Pro Pro Ala Glu Leu Glu Ser val Leu
 10 470
 Leu Gln His Pro Asn Ile Phe Asp Ala Gly
 480
 Val Ala Gly Val Pro Asp Pro Ile Ala Gly
 15 490
 Glu Leu Pro Gly Ala Val Val Val Leu Glu
 500
 20 Lys Gly Lys Ser Met Thr Glu Lys Glu Val
 510
 Met Asp Tyr Val Ala Ser Gln Val Ser Asn
 25 520
 Ala Lys Arg Leu Arg Gly Gly Val Arg Phe
 530
 Val Asp Glu Val Pro Lys Gly Leu Thr Gly
 30 540
 Lys Ile Asp Gly Lys Ala Ile Arg Glu Ile
 35
 Leu Lys Lys Pro Val Ala Lys Met

3. Gène de luciférase selon la revendication 1 ou 2 qui est représenté par la séquence nucléotidique
 40 présentée ci-dessous:

30
 45 ATG GAA AAC ATG GAG AAC GAT GAA AAT ATT
 60
 GTG TAT GGT CCT GAA CCA TTT TAC CCT ATT
 90
 50 GAA GAG GGA TCT GCT GGA GCA CAA TTG CGC
 55

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		120
	AAG TAT ATG GAT CGA TAT GCA AAA CTT GGA	
5		150
	GCA ATT GCT TTT ACT AAC GCA CTT ACC GGT	
		180
10	GTC GAT TAT ACG TAC GCC GAA TAC TTA GAA	
		210
	AAA TCA TGC TGT CTA GGA GAG GCT TTA AAG	
		240
15	AAT TAT GGT TTG GTT GTT GAT GGA AGA ATT	
		270
	GCG TTA TGC AGT GAA AAC TGT GAA GAA TTC	
20		300
	TTT ATT CCT GTA TTA GCC GGT TTA TTT ATA	
		330
25	GGT GTC GGT GTG GCT CCA ACT AAT GAG ATT	
		360
	TAC ACT CTA CGT GAA TTG GTT CAC AGT TTA	
		390
30	GGC ATC TCT AAG CCA ACA ATT GTA TTT AGT	
		420
	TCT AAA AAA GGA TTA GAT AAA GTT ATA ACT	
35		450
	GTA CAA AAA ACG GTA ACT GCT ATT AAA ACC	
		480
40	ATT GTT ATA TTG GAC AGC AAA GTG GAT TAT	
		510
	AGA GGT TAT CAA TCC ATG GAC AAC TTT ATT	
		540
45	AAA AAA AAC ACT CCA CAA GGT TTC AAA GGA	
		570
	TCA AGT TTT AAA ACT GTA GAA GTT AAC CGC	
50		600
	AAA GAA CAA GTT GCT CTT ATA ATG AAC TCT	

55

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		630
	TCG GGT TCA ACC GGT TTG CCA AAA GGT GTG	
5		660
	CAA CTT ACT CAT GAA AAT GCA GTC ACT AGA	
		690
10	TTT TCT CAC GCT AGA GAT CCA ATT TAT GGA	
		720
	AAC CAA GTT TCA CCA GGC ACG GCT ATT TTA	
		750
15	ACT GTA GTA CCA TTC CAT CAT GGT TTT GGT	
		780
	ATG TTT ACT ACT TTA GGC TAT CTA ACT TGT	
20		810
	GGT TTT CGT ATT GTC ATG TTA ACG AAA TTT	
		840
	GAC GAA GAG ACT TTT TTA AAA ACA CTG CAA	
25		870
	GAT TAC AAA TGT TCA AGC GTT ATT CTT GTA	
		900
30	CCG ACT TTG TTT GCA ATT CTT AAT AGA AGT	
		930
	GAA TTA CTC GAT AAA TAT GAT TTA TCA AAT	
35		960
	TTA GTT GAA ATT GCA TCT GGC GGA GCA CCT	
		990
40	TTA TCT AAA GAA ATT GGT GAA GCT GTT GCT	
		1020
	AGA CGT TTT AAT TTA CCG GGT GTT CGT CAA	
		1050
45	GGC TAT GGT TTA ACA GAA ACA ACC TCT GCA	
		1080
	ATT ATT ATC ACA CCG GAA GGC GAT GAT AAA	
50		1110
	CCA GGT GCT TCT GGC AAA GTT GTG CAA TTA	

55

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1140
 TTT AAA GCA AAA GTT ATC GAT CTT GAT ACT
 5 1170
 AAA AAA ACT TTG GGC CCG AAC AGA CGT GGA
 1200
 GAA GTT TGT GTA AAG GGT CCT ATG CTT ATG
 10 1230
 AAA GGT TAT GTA GAT AAT CCA GAA GCA ACA
 1260
 15 AGA GAA ATC ATA GAT GAA GAA GGT TGG TTG
 1290
 CAC ACA GGA GAT ATT GGG TAT TAC GAT GAA
 20 1320
 GAA AAA CAT TTC TTT ATC GTG GAT CGT TTG
 1350
 AAG TCT TTA ATC AAA TAC AAA GGA TAT CAA
 25 1380
 GTA CCA CCT GCT GAA TTA GAA TCT GTT CTT
 1410
 30 TTG CAA CAT CCA AAT ATT TTT GAT GCC GGC
 1440
 GTT GCT GGC GTT CCA GAT CCT ATA GCT GGT
 35 1470
 GAG CTT CCG GGA GCT GTT GTT GTA CTT GAA
 1500
 AAA GGA AAA TCT ATG ACT GAA AAA GAA GTA
 40 1530
 ATG GAT TAC GTT GCT AGT CAA GTT TCA AAT
 1560
 45 GCA AAA CGT TTG CGT GGT GGT GTC CGT TTT
 1590
 GTG GAC GAA GTA CCT AAA GGT CTC ACT GGT
 50 1620
 AAA ATT GAC GGT AAA GCA ATT AGA GAA ATA
 55
 CTG AAG AAA CCA GTT GCT AAG ATG

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4. Nouvel ADN recombiné comprenant un ADN vecteur dans lequel est inséré un gène codant la luciférase dérivée de Luciola lateralis selon les revendications 1 à 3.
5. Nouvel ADN recombiné selon la revendication 4, dans lequel ledit ADN vecteur est le plasmide pUC119.
6. Procédé de production d'une luciférase qui comprend la culture dans un milieu d'un micro-organisme appartenant au genre *Escherichia* et portant un ADN recombiné selon la revendication 4 obtenu par insertion d'un gène codant la luciférase dérivée de Luciola lateralis dans un ADN vecteur et l'obtention de ladite luciférase à partir de la culture.
7. Procédé de production d'une luciférase selon la revendication 6, dans lequel ledit ADN vecteur est le plasmide pUC119.

FIG. 1

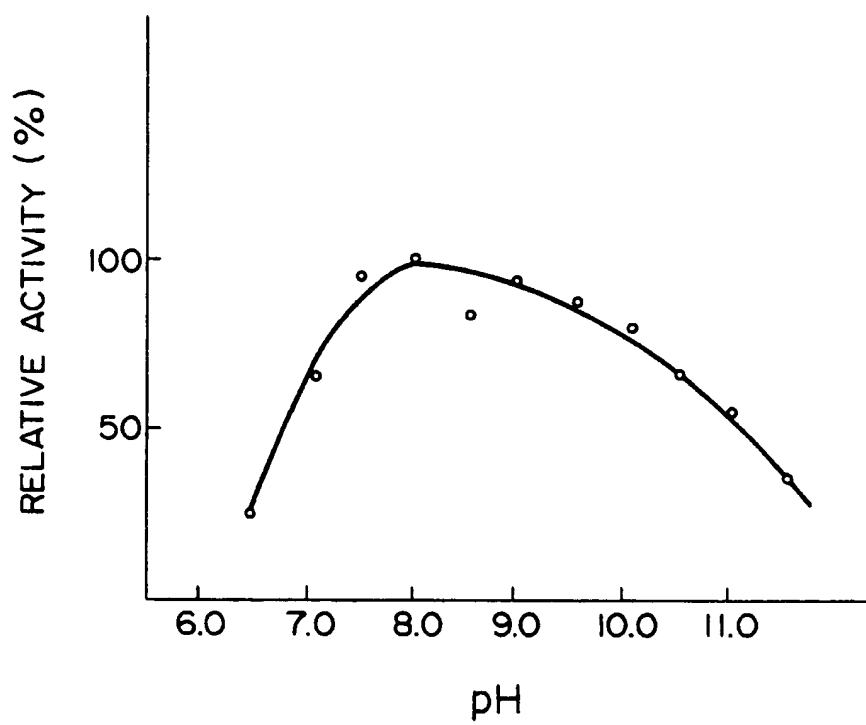


FIG. 2

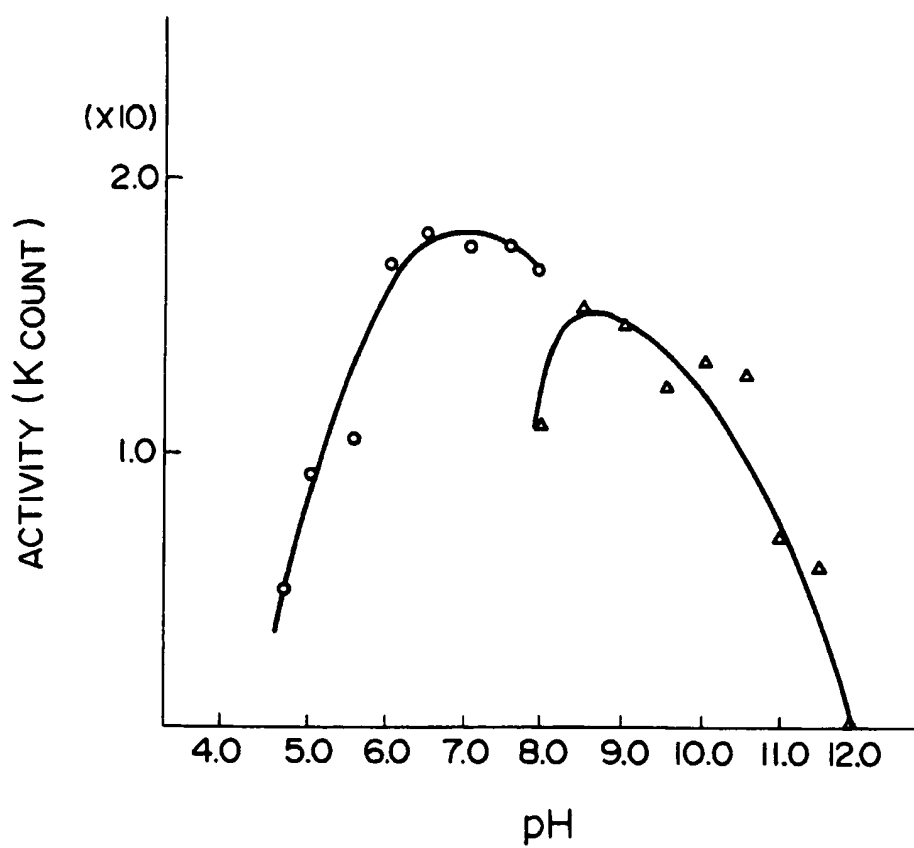


FIG. 3

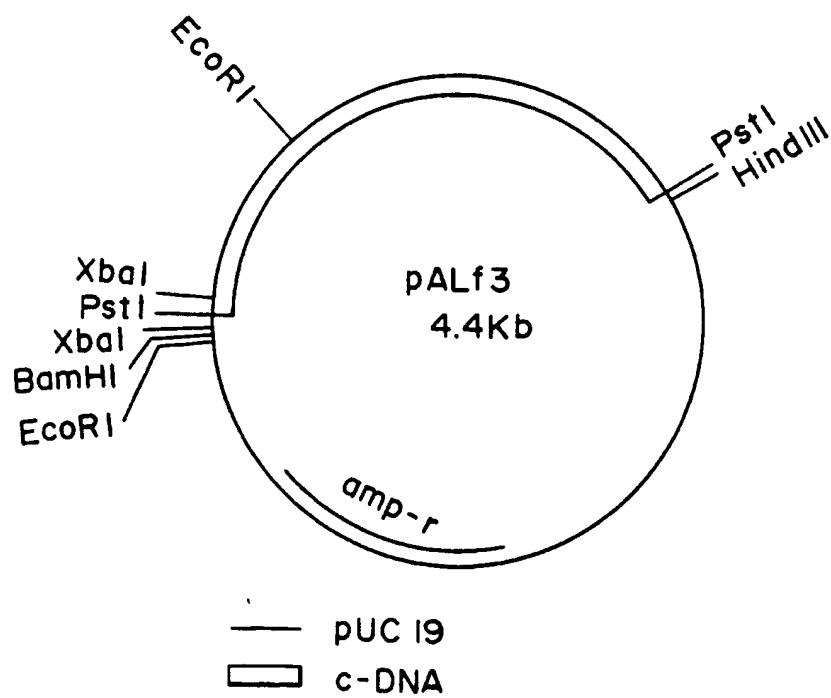


FIG. 4

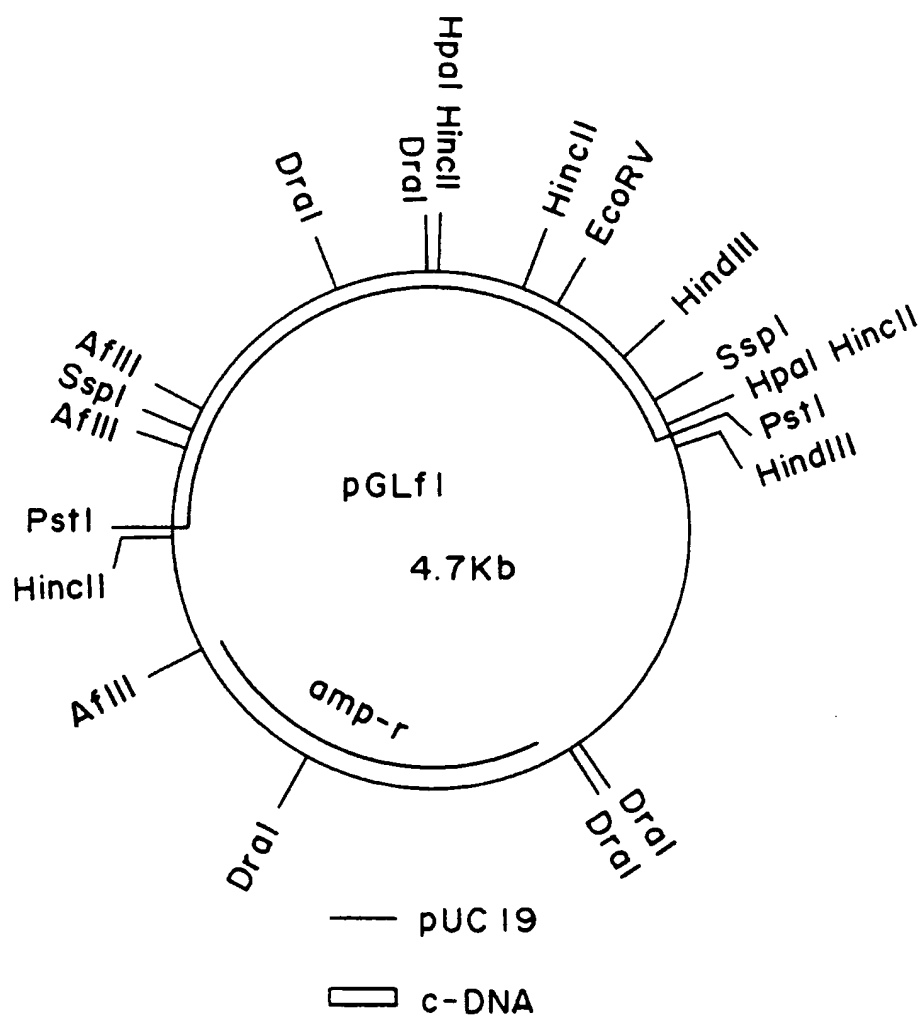


FIG. 5

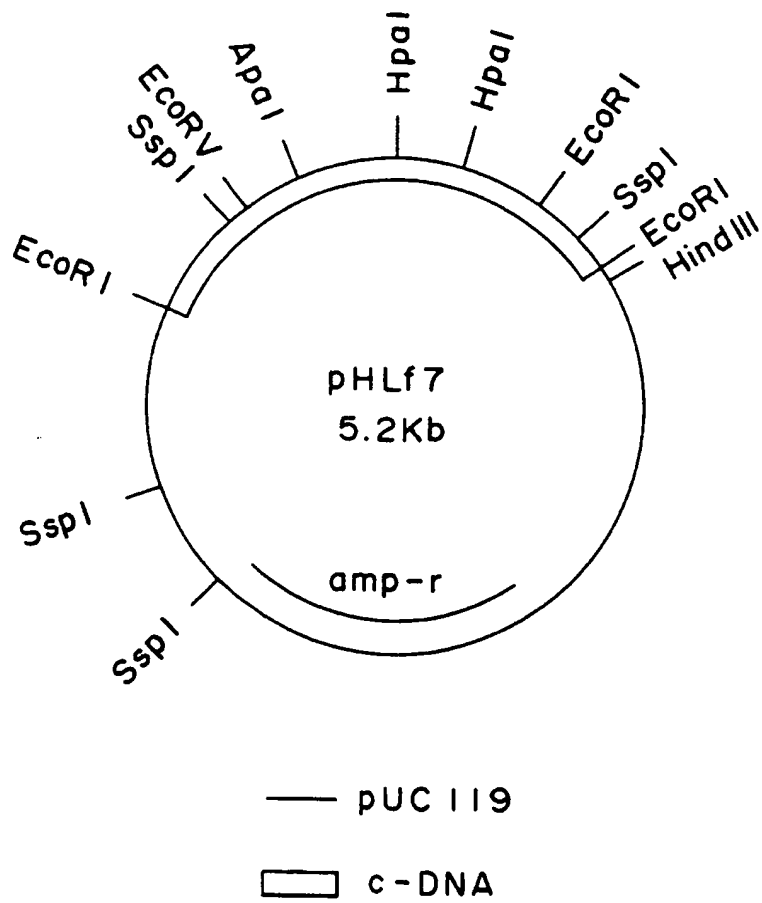


FIG. 6A

```

ATG GAA AAC ATG GAG AAC GAT GAA AAT ATT 30
GTG TAT GGT CCT GAA CCA TTT TAC CCT ATT 60
GAA GAG GGA TCT GCT GGA GCA CAA TTG CGC 90
AAG TAT ATG GAT CGA TAT GCA AAA CTT GGA 120
GCA ATT GCT TTT ACT AAC GCA CTT ACC GGT 150
GTC GAT TAT ACG TAC GCC GAA TAC TTA GAA 180
AAA TCA TGC TGT CTA GGA GAG GCT TTA AAG 210
AAT TAT GGT TTG GTT GTT GAT GGA AGA ATT 240
GCG TTA TGC AGT GAA AAC TGT GAA GAA TTC 270
TTT ATT CCT GTA TTA GCC GGT TTA TTT ATA 300
GGT GTC GGT GTG GCT CCA ACT AAT GAG ATT 330
TAC ACT CTA CGT GAA TTG GTT CAC AGT TTA 360
GGC ATC TCT AAG CCA ACA ATT GTA TTT AGT 390
TCT AAA AAA GGA TTA GAT AAA GTT ATA ACT 420
GTA CAA AAA ACG GTA ACT GCT ATT AAA ACC 450
ATT GTT ATA TTG GAC AGC AAA GTG GAT TAT 480
AGA GGT TAT CAA TCC ATG GAC AAC TTT ATT 510
AAA AAA AAC ACT CCA CAA GGT TTC AAA GGA 540
TCA AGT TTT AAA ACT GTA GAA GTT AAC CGC 570

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FIG. 6B

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AAA GAA CAA GTT GCT CTT ATA ATG AAC TCT 600
TCG GGT TCA ACC GGT TTG CCA AAA GGT GTG 630
CAA CTT ACT CAT GAA AAT GCA GTC ACT AGA 660
TTT TCT CAC GCT, AGA GAT CCA ATT TAT GGA 690
AAC CAA GTT TCA CCA GGC ACG GCT ATT TTA 720
ACT GTA GTA CCA TTC CAT CAT GGT TTT GGT 750
ATG TTT ACT ACT TTA GGC TAT CTA ACT TGT 780
GGT TTT CGT ATT GTC ATG TTA ACG AAA TTT 810
GAC GAA GAG ACT TTT TTA AAA ACA CTG CAA 840
GAT TAC AAA TGT TCA AGC GTT ATT CTT GTA 870
CCG ACT TTG TTT GCA ATT CTT AAT AGA AGT 900
GAA TTA CTC GAT AAA TAT GAT TTA TCA AAT 930
TTA GTT GAA ATT GCA TCT GGC GGA GCA CCT 960
TTA TCT AAA GAA ATT GGT GAA GCT GTT GCT 990
AGA CGT TTT AAT TTA CCG GGT GTT CGT CAA 1020
GGC TAT GGT TTA ACA GAA ACA ACC TCT GCA 1050
ATT ATT ATC ACA CCG GAA GGC GAT GAT AAA 1080
CCA GGT GCT TCT GGC AAA GTT GTG CCA TTA 1110
TTT AAA GCA AAA GTT ATC GAT CTT GAT ACT 1140

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FIG. 6C

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AAA AAA ACT TTG GGC CCG AAC AGA CGT GGA 1170
GAA GTT TGT GTA AAG GGT CCT ATG CTT ATG 1200
AAA GGT TAT GTA GAT AAT CCA GAA GCA ACA 1230
AGA GAA ATC ATA GAT GAA GAA GGT TGG TTG 1260
CAC ACA GGA GAT ATT GGG TAT TAC GAT GAA 1290
GAA AAA CAT TTC TTT ATC GTG GAT CGT TTG 1320
AAG TCT TTA ATC AAA TAC AAA GGA TAT CAA 1350
GTA CCA CCT GCT GAA TTA GAA TCT GTT CTT 1380
TTG CAA CAT CCA AAT ATT TTT GAT GCC GGC 1410
GTT GCT GGC GTT CCA GAT CCT ATA GCT GGT 1440
GAG CTT CCG GGA GCT GTT GTT GTA CTT GAA 1470
AAA GGA AAA TCT ATG ACT GAA AAA GAA GTA 1500
ATG GAT TAC GTT GCT AGT CAA GTT TCA AAT 1530
GCA AAA CGT TTG CGT GGT GGT GTC CGT TTT 1560
GTG GAC GAA GTA CCT AAA GGT CTC ACT GGT 1590
AAA ATT GAC GGT AAA GCA ATT AGA GAA ATA 1620
CTG AAG AAA CCA GTT GCT AAG ATG

```

FIG. 7A

```

Met Glu Asn Met Glu Asn Asp Glu Asn Ile 10
Val Tyr Gly Pro Glu Pro Phe Tyr Pro Ile 20
Glu Glu Gly Ser Ala Gly Ala Gln Leu Arg 30
Lys Tyr Met Asp Arg Tyr Ala Lys Leu Gly 40
Ala Ile Ala Phe Thr Asn Ala Leu Thr Gly 50
Val Asp Tyr Thr Tyr Ala Glu Tyr Leu Glu 60
Lys Ser Cys Cys Leu Gly Glu Ala Leu Lys 70
Asn Tyr Gly Leu Val Val Asp Gly Arg Ile 80
Ala Leu Cys Ser Glu Asn Cys Glu Glu Phe 90
Phe Ile Pro Val Leu Ala Gly Leu Phe Ile 100
Gly Val Gly Val Ala Pro Thr Asn Glu Ile 110
Tyr Thr Leu Arg Glu Leu Val His Ser Leu 120
Gly Ile Ser Lys Pro Thr Ile Val Phe Ser 130
Ser Lys Lys Gly Leu Asp Lys Val Ile Thr 140
Val Gln Lys Thr Val Thr Ala Ile Lys Thr 150
Ile Val Ile Leu Asp Ser Lys Val Asp Tyr 160
Arg Gly Tyr Gln Ser Met Asp Asn Phe Ile 170
Lys Lys Asn Thr Pro Gln Gly Phe Lys Gly 180
Ser Ser Phe Lys Thr Val Glu Val Asn Arg 190

```

FIG. 7B

Lys	Glu	Gln	Val	Ala	Leu	Ile	Met	Asn	Ser	200
Ser	Gly	Ser	Thr	Gly	Leu	Pro	Lys	Gly	Val	210
Gln	Leu	Thr	His	Glu	Asn	Ala	Val	Thr	Arg	220
Phe	Ser	His	Ala	Arg	Asp	Pro	Ile	Tyr	Gly	230
Asn	Gln	Val	Ser	Pro	Gly	Thr	Ala	Ile	Leu	240
Thr	Val	Val	Pro	Phe	His	His	Gly	Phe	Gly	250
Met	Phe	Thr	Thr	Leu	Gly	Tyr	Leu	Thr	Cys	260
Gly	Phe	Arg	Ile	Val	Met	Leu	Thr	Lys	Phe	270
Asp	Glu	Glu	Thr	Phe	Leu	Lys	Thr	Leu	Gln	280
Asp	Tyr	Lys	Cys	Ser	Ser	Val	Ile	Leu	Val	290
Pro	Thr	Leu	Phe	Ala	Ile	Leu	Asn	Arg	Ser	300
Glu	Leu	Leu	Asp	Lys	Tyr	Asp	Leu	Ser	Asn	310
Leu	Val	Glu	Ile	Ala	Ser	Gly	Gly	Ala	Pro	320
Leu	Ser	Lys	Glu	Ile	Gly	Glu	Ala	Val	Ala	330
Arg	Arg	Phe	Asn	Leu	Pro	Gly	Val	Arg	Gln	340
Gly	Tyr	Gly	Leu	Thr	Glu	Thr	Thr	Ser	Ala	350
Ile	Ile	Ile	Thr	Pro	Glu	Gly	Asp	Asp	Lys	360
Pro	Gly	Ala	Ser	Gly	Lys	Val	Val	Pro	Leu	370
Phe	Lys	Ala	Lys	Val	Ile	Asp	Leu	Asp	Thr	380

FIG. 7C

```

Lys Lys Thr Leu Gly Pro Asn Arg Arg Gly 390
Glu Val Cys Val Lys Gly Pro Met Leu Met 400
Lys Gly Tyr Val Asp Asn Pro Glu Ala Thr 410
Arg Glu Ile Ile Asp Glu Glu Gly Trp Leu 420
His Thr Gly Asp Ile Gly Tyr Tyr Asp Glu 430
Glu Lys His Phe Phe Ile Val Asp Arg Leu 440
Lys Ser Leu Ile Lys Tyr Lys Gly Tyr Gln 450
Val Pro Pro Ala Glu Leu Glu Ser Val Leu 460
Leu Gln His Pro Asn Ile Phe Asp Ala Gly 470
Val Ala Gly Val Pro Asp Pro Ile Ala Gly 480
Glu Leu Pro Gly Ala Val Val Val Leu Glu 490
Lys Gly Lys Ser Met Thr Glu Lys Glu Val 500
Met Asp Tyr Val Ala Ser Gln Val Ser Asn 510
Ala Lys Arg Leu Arg Gly Gly Val Arg Phe 520
Val Asp Glu Val Pro Lys Gly Leu Thr Gly 530
Lys Ile Asp Gly Lys Ala Ile Arg Glu Ile 540
Leu Lys Lys Pro Val Ala Lys Met

```

FIG. 8A

ATG	GAA	AAC	ATG	GAG	AAC	GAT	GAA	AAT	ATT	10
Met	Glu	Asn	Met	Glu	Asn	Asp	Glu	Asn	Ile	
GTG	TAT	GGT	CCT	GAA	CCA	TTT	TAC	CCT	ATT	20
Val	Tyr	Gly	Pro	Glu	Pro	Phe	Tyr	Pro	Ile	
GAA	GAG	GGA	TCT	GCT	GGA	GCA	CAA	TTG	CGC	30
Glu	Glu	Gly	Ser	Ala	Gly	Ala	Gln	Leu	Arg	
AAG	TAT	ATG	GAT	CGA	TAT	GCA	AAA	CTT	GGA	40
Lys	Tyr	Met	Asp	Arg	Tyr	Ala	Lys	Leu	Gly	
GCA	ATT	GCT	TTT	ACT	AAC	GCA	CTT	ACC	GGT	50
Ala	Ile	Ala	Phe	Thr	Asn	Ala	Leu	Thr	Gly	
GTC	GAT	TAT	ACG	TAC	GCC	GAA	TAC	TTA	GAA	60
Val	Asp	Tyr	Thr	Tyr	Ala	Glu	Tyr	Leu	Glu	
AAA	TCA	TGC	TGT	CTA	GGA	GAG	GCT	TTA	AAG	70
Lys	Ser	Cys	Cys	Leu	Gly	Glu	Ala	Leu	Lys	
AAT	TAT	GGT	TTG	GTT	GTT	GAT	GGA	AGA	ATT	80
Asn	Tyr	Gly	Leu	Val	Val	Asp	Gly	Arg	Ile	
GCG	TTA	TGC	AGT	GAA	AAC	TGT	GAA	GAA	TTC	90
Ala	Leu	Cys	Ser	Glu	Asn	Cys	Glu	Glu	Phe	
TTT	ATT	CCT	GTA	TTA	GCC	GGT	TTA	TTT	ATA	100
Phe	Ile	Pro	Val	Leu	Ala	Gly	Leu	Phe	Ile	
GGT	GTC	GGT	GTG	GCT	CCA	ACT	AAT	GAG	ATT	110
Gly	Val	Gly	Val	Ala	Pro	Thr	Asn	Glu	Ile	
TAC	ACT	CTA	CGT	GAA	TTG	GTT	CAC	AGT	TTA	120
Tyr	Thr	Leu	Arg	Glu	Leu	Val	His	Ser	Leu	

FIG. 8B

GGC	ATC	TCT	AAG	CCA	ACA	ATT	GTA	TTT	AGT	130
Gly	Ile	Ser	Lys	Pro	Thr	Ile	Val	Phe	Ser	
TCT	AAA	AAA	GGA	TTA	GAT	AAA	GTT	ATA	ACT	140
Ser	Lys	Lys	Gly	Leu	Asp	Lys	Val	Ile	Thr	
GTA	CAA	AAA	ACG	GTA	ACT	GCT	ATT	AAA	ACC	150
Val	Gln	Lys	Thr	Val	Thr	Ala	Ile	Lys	Thr	
ATT	GTT	ATA	TTG	GAC	AGC	AAA	GTG	GAT	TAT	160
Ile	Val	Ile	Leu	Asp	Ser	Lys	Val	Asp	Tyr	
AGA	GGT	TAT	CAA	TCC	ATG	GAC	AAC	TTT	ATT	170
Arg	Gly	Tyr	Gln	Ser	Met	Asp	Asn	Phe	Ile	
AAA	AAA	AAC	ACT	CCA	CAA	GGT	TTC	AAA	GGA	180
Lys	Lys	Asn	Thr	Pro	Gln	Gly	Phe	Lys	Gly	
TCA	AGT	TTT	AAA	ACT	GTA	GAA	GTT	AAC	CGC	190
Ser	Ser	Phe	Lys	Thr	Val	Glu	Val	Asn	Arg	
AAA	GAA	CAA	GTT	GCT	CTT	ATA	ATG	AAC	TCT	200
Lys	Glu	Gln	Val	Ala	Leu	Ile	Met	Asn	Ser	
TCG	GGT	TCA	ACC	GGT	TTG	CCA	AAA	GGT	GTG	210
Ser	Gly	Ser	Thr	Gly	Leu	Pro	Lys	Gly	Val	
CAA	CTT	ACT	CAT	GAA	AAT	GCA	GTC	ACT	AGA	220
Gln	Leu	Thr	His	Glu	Asn	Ala	Val	Thr	Arg	
TTT	TCT	CAC	GCT	AGA	GAT	CCA	ATT	TAT	GGA	230
Phe	Ser	His	Ala	Arg	Asp	Pro	Ile	Tyr	Gly	
AAC	CAA	GTT	TCA	CCA	GGC	ACG	GCT	ATT	TTA	240
Asn	Gln	Val	Ser	Pro	Gly	Thr	Ala	Ile	Leu	

FIG. 8C

ACT	GTA	GTA	CCA	TTC	CAT	CAT	GGT	TTT	GGT	250
Thr	Val	Val	Pro	Phe	His	His	Gly	Phe	Gly	
ATG	TTT	ACT	ACT	TTA	GGC	TAT	CTA	ACT	TGT	260
Met	Phe	Thr	Thr	Leu	Gly	Tyr	Leu	Thr	Cys	
GGT	TTT	CGT	ATT	GTC	ATG	TTA	ACG	AAA	TTT	270
Gly	Phe	Arg	Ile	Val	Met	Leu	Thr	Lys	Phe	
GAC	GAA	GAG	ACT	TTT	TTA	AAA	ACA	CTG	CAA	280
Asp	Glu	Glu	Thr	Phe	Leu	Lys	Thr	Leu	Gln	
GAT	TAC	AAA	TGT	TCA	AGC	GTT	ATT	CTT	GTA	290
Asp	Tyr	Lys	Cys	Ser	Ser	Val	Ile	Leu	Val	
CCG	ACT	TTG	TTT	GCA	ATT	CTT	AAT	AGA	AGT	300
Pro	Thr	Leu	Phe	Ala	Ile	Leu	Asn	Arg	Ser	
GAA	TTA	CTC	GAT	AAA	TAT	GAT	TTA	TCA	AAT	310
Glu	Leu	Leu	Asp	Lys	Tyr	Asp	Leu	Ser	Asn	
TTA	GTT	GAA	ATT	GCA	TCT	GGC	GGA	GCA	CCT	320
Leu	Val	Glu	Ile	Ala	Ser	Gly	Gly	Ala	Pro	
TTA	TCT	AAA	GAA	ATT	GGT	GAA	GCT	GTT	GCT	330
Leu	Ser	Lys	Glu	Ile	Gly	Glu	Ala	Val	Ala	
AGA	CGT	TTT	AAT	TTA	CCG	GGT	GTT	CGT	CAA	340
Arg	Arg	Phe	Asn	Leu	Pro	Gly	Val	Arg	Gln	
GGC	TAT	GGT	TTA	ACA	GAA	ACA	ACC	TCT	GCA	350
Gly	Tyr	Gly	Leu	Thr	Glu	Thr	Thr	Ser	Ala	
ATT	ATT	ATC	ACA	CCG	GAA	GGC	GAT	GAT	AAA	360
Ile	Ile	Ile	Thr	Pro	Glu	Gly	Asp	Asp	Lys	

FIG. 8D

CCA	GGT	GCT	TCT	GGC	AAA	GTT	GTG	CCA	TTA	370
Pro	Gly	Ala	Ser	Gly	Lys	Val	Val	Pro	Leu	
TTT	AAA	GCA	AAA	GTT	ATC	GAT	CTT	GAT	ACT	380
Phe	Lys	Ala	Lys	Val	Ile	Asp	Leu	Asp	Thr	
AAA	AAA	ACT	TTG	GGC	CCG	AAC	AGA	CGT	GGA	390
Lys	Lys	Thr	Leu	Gly	Pro	Asn	Arg	Arg	Gly	
GAA	GTT	TGT	GTA	AAG	GGT	CCT	ATG	CTT	ATG	400
Glu	Val	Cys	Val	Lys	Gly	Pro	Met	Leu	Met	
AAA	GGT	TAT	GTA	GAT	AAT	CCA	GAA	GCA	ACA	410
Lys	Gly	Tyr	Val	Asp	Asn	Pro	Glu	Ala	Thr	
AGA	GAA	ATC	ATA	GAT	GAA	GAA	GGT	TGG	TTG	420
Arg	Glu	Ile	Ile	Asp	Glu	Glu	Gly	Trp	Leu	
CAC	ACA	GGA	GAT	ATT	GGG	TAT	TAC	GAT	GAA	430
His	Thr	Gly	Asp	Ile	Gly	Tyr	Tyr	Asp	Glu	
GAA	AAA	CAT	TTC	TTT	ATC	GTG	GAT	CGT	TTG	440
Glu	Lys	His	Phe	Phe	Ile	Val	Asp	Arg	Leu	
AAG	TCT	TTA	ATC	AAA	TAC	AAA	GGA	TAT	CAA	450
Lys	Ser	Leu	Ile	Lys	Tyr	Lys	Gly	Tyr	Gln	
GTA	CCA	CCT	GCT	GAA	TTA	GAA	TCT	GTT	CTT	460
Val	Pro	Pro	Ala	Glu	Leu	Glu	Ser	Val	Leu	
TTG	CAA	CAT	CCA	AAT	ATT	TTT	GAT	GCC	GGC	470
Leu	Gln	His	Pro	Asn	Ile	Phe	Asp	Ala	Gly	
GTT	GCT	GGC	GTT	CCA	GAT	CCT	ATA	GCT	GGT	480
Val	Ala	Gly	Val	Pro	Asp	Pro	Ile	Ala	Gly	

FIG. 8E

GAG	CTT	CCG	GGA	GCT	GTT	GTT	GTA	CTT	GAA	490
Glu	Leu	Pro	Gly	Ala	Val	Val	Val	Leu	Glu	
AAA	GGA	AAA	TCT	ATG	ACT	GAA	AAA	GAA	GTA	500
Lys	Gly	Lys	Ser	Met	Thr	Glu	Lys	Glu	Val	
ATG	GAT	TAC	GTT	GCT	AGT	CAA	GTT	TCA	AAT	510
Met	Asp	Tyr	Val	Ala	Ser	Gln	Val	Ser	Asn	
GCA	AAA	CGT	TTG	CGT	GGT	GGT	GTC	CGT	TTT	520
Ala	Lys	Arg	Leu	Arg	Gly	Gly	Val	Arg	Phe	
GTG	GAC	GAA	GTA	CCT	AAA	GGT	CTC	ACT	GGT	530
Val	Asp	Glu	Val	Pro	Lys	Gly	Leu	Thr	Gly	
AAA	ATT	GAC	GGT	AAA	GCA	ATT	AGA	GAA	ATA	540
Lys	Ile	Asp	Gly	Lys	Ala	Ile	Arg	Glu	Ile	
CTG	AAG	AAA	CCA	GTT	GCT	AAG	ATG			
Leu	Lys	Lys	Pro	Val	Ala	Lys	Met			